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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Title:
Perfect score:
                                                                                                                                                                                                                                                                                                           Minimum DB seq length: 0
Maximum DB seq length: 2000000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OM protein - protein search, using sw model
                                                                                                                                                                                                                                 Database
                                                                                                                                                                                                                                                                                                                                                   Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                               Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                   Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence:
                                                                                                                                                                                       SPTREMBL 25:*

1: sp_archea:*

2: sp_bacteria:*

3: sp_fungi:*
                                                                                                                                                                                                                                                                                                                                                                                                   BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                                              US-10-612-885A-1
117
1 QRVEILEGRTECVL
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6:
7:
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111:
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113:
115:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                   sp_human:*
sp_invertebrate:*
sp_mammal:*
sp_mhc:*
sp_lant:*
sp_rodent:*
sp_virus:*
sp_vartebrate:*
sp_unclassified:*
sp_rvirus:*
sp_bacteriap:*
sp_archeap:*
                                                                                                            sp_organelle:*
sp_phage:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6, 2004, 12:41:36; Search time 39 Seconds (without alignments) 186.075 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                      1017041
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

16	15	14	13	12	11	10	9	80	7	o,	ហ	4	ω	2	_	Result No.
47	47	47	48	48	48	49	49	52	56	104	104	104	104	107	109	Score
40.2	40.2	40.2	41.0	41.0	41.0	41.9	41.9	44.4	47.9	88.9	88.9	88.9	88.9	91.5	93.2	Query Match
640	539	539	1028	1028	428	1028	826	1028	176	418	387	229	229	509	316	Length
10	10	10	11	11	10	片	N	4	17	σ	σ	σ	σ	σ	11	BB
Q9ASX3	Q7XDN4	Q94LS6	Q8C6X1	8EMD6	Q9LPY7	P97528	Q9AKX8	Q9UQ52	Q8TYF5	Q95N14	Q95N13	Q27950	Q28206	Q9MYZ9	035545	ID
Q9asx3 arabidopsis	Q7xdn4 oryza sativ	Q941s6 oryza sativ		Q9jmb8 mus musculu	Q9lpy7 arabidopsis	P97528 rattus norv	Q9akx8 legionella	Q9uq52 homo sapien	Q8tyf5 methanopyru	Q95n14 ovis aries	Q95n13 ovis aries	Q27950 bos indicus	Q28206 bos taurus	Q9myz9 sus scrofa	035545 rattus norv	Description

45	44	43	42	41	40	39	38	37	36	35	34	ပ္သ	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17
44	44	44	44	44.5			4 5	4 5	45	45	45	45	45	45	45	4 5	4 5	45.5	46	46	46	46	46	46	47	47	47	47
37.6	37.6	7	7	38.0	æ	æ	38.5	ω	α	α	8	æ	8	œ	æ	æ	æ	œ	9	9	9	9	9	9	0	0	40.2	40.2
2023	606	195	47	810	693	691	1056	968	773	567	567	479	446	316	81	76	76	95	1283	781	700	700	637	288	1873	1226	644	644
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091632	Q9VWF6	Q8Z0G3	Q8XFL2	Q9ZWI3	096680	Q9VML1	057161	Q7YZ47	Q8A0C0	Q9VCA3	Q95R48	Q9U7PS	Q94CT7	Q912J3	Q30747	Q30567	Q30734	Q8TF83	Q8TA82	Q82857	Q8C0B3	Q9CXF5	Q86GJ9	Q98214	Q9FGI1	Q7UL12	Q9FNK3	Q8LDV1
O91632 cherry gree	irosophila	Q8z0g3 anabaena sp	Q8xfl2 salmonella		O96680 drosophila	Q9vml1 drosophila		Q7yz47 cryptospori	Q8a0c0 bacteroides	Ω,	Q95r48 drosophila	Q9u7p5 eufolliculi		Q9i2j3 pseudomonas	3	Q30567 macaca fasc	macaca	Q8tf83 agrocybe ae	a	7	Q8c0b3 mus musculu	Q9cxf5 mus musculu	dict	4	Q9fgil arabidopsis	Q7ul12 rhodopirell	Q9fnk3 arabidopsis	

ALIGNMENTS

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RESULT 3
Q282DC Q282
AC Q282
AC Q282
AC Q282
DT 01-N
DT 01-N
DT 01-O
DE EFFOR
GN EPOR
OS BOS
OC ENTER
OC BOWING
OC Mammi
OC SEQU
RC TISS
RA Sulli
RL Submi
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REMBL, AF274305; AAF77065.1; -.

REMBL, AF274305; AERN.

RGO; GO:0016020; C:membrane; IBA.

RGO; GO:0016020; C:membrane; IBA.

RGO; GO:0004872; F:receptor activity; IEA.

RGO; GO:0004872; F:receptor activity; IEA.

RGO; GO:0006118; P:electron transport; IEA.

RGO; GO:0006118; P:electron transport; IEA.

RINterPro; IPR002966; CRIA.

RINterPro; IPR003961; FN_III.

RINterPro; IPR003961; FN_III-1ike.

RINterPro; IPR003952; Hemtopoptn L.FI.

DR InterPro; IPR003528; Hemtopoptn L.FI.

SMART; SM00060; FN3; 1.

SMART; SM00060; FN3; 1.

ENART; SM00060; FN3; 1.
                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 21
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                                                                                                                                                     Q28206;
Q28206;
01-NOV-1996
01-NOV-1996
01-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q9MYZ9
Q9MYZ9;
01-OCT-2000
 TISSUE=Bone marrow;
Suliman H.B., Feldman
Submitted (JUN-1996) t
                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea
Bovidae; Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                                                                          Receptor.
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Porcine Erythropoietin Receptor: Molecular Cloning and Embryonic and Fetal Liver.; Domest. Anim. Endocrinol. 0:0-0(2000). EMBL; AF274305; AAF77065.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                 PROSITE; PS01352; HEMATOPO REC L F1; PROSITE; PS00559; MOLYBDOPTERIN_EUK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pearson P.L., Smith T.P.L., So Christenson R.K., Vallet J.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa;
Mammalia; Eutheria;
                                     SEQUENCE FROM N.A.
                                                                                                                Bos taurus (Bovine).
                                                                                                                                        Erythropoietin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=9823;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sus scrofa (Pig).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Erythropoietin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-OCT-2003
                                                             NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TISSUE=Liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                     QRVEILEGRTECVLSNLRGRTRY 23
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                                                                                                                                                                                                                                                                         QRVEILDGRTECVLSNLRGGTRY 217
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                                                                                                                                                                                                                                                                                                                                                                            509
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                        (TYENMBLrel. 01, Created)
(TYENMBLrel. 01, Last sequence up
(TYENMBLrel. 25, Last annotation
in receptor (Fragment).
                                                                                                                                                                                                          PRELIMINARY;
                                                                                                                                                                                                                                                                                                                         Conservative
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В
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 .F.,
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Pred. No. 2.5e
1; Mismatches
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Last sequence update)
Last annotation update)
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 Majiwa P.A.O., Logan-Henfrey EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7.
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                                                                                                                                                                                                                                                                                                                                                                          35B565D07C6BCD8A CRC64;
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on update)
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01-NOV-1996
01-NOV-1996
01-OCT-2003
                                                                                                                                                                 Receptor.
NON_TER
NON_TER
SEQUENCE
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NON_TER
NON_TER
                                                                                                                                                                                                                                                SMART; SM
PROSITE;
                                                                                                                                                                                                                                                                                                                                             GO; GO:0016020; C:membrane; IEA.
GO; GO:0004956; F:hematopoletin/Interferon-class
GO; GO:0004872; F:receptor activity; IEA.
InterPro; IPR002996; CR1A.
InterPro; IPR003961; FN_III.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro; IPR003961; FN_III.
InterPro; IPR003957; FN_III-like.
InterPro; IPR003952; Hemtopoptn_L_F1.
Pfam; PF00041; En3; 1.
SMART; SM00060; FN3; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Suliman H.B., Feldman B.F., Submitted (JUN-1996) to the EMBL; U61398; AAB03870.1; -. HSSP; P19235; 1EBA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A. TISSUE=Bone marrow;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa;
Mammalia; Eutheria;
Bovidae; Bovinae; Bo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Erythropoietin receptor
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GO; GO:0004896; F:hematopoietin/interferon-class (D200-domain.
GO; GO:0004872; F:receptor activity; IEA.
InterPro; IPR002996; CR1A.
                                                                                                                                                                                                                                                                                          Pfam; PF00041; fn3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bos indicus (Zebu).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Q27950
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                                                                                                                                                                                                                                                                                                               InterPro;
                                                                                                                                                                                                                                                                                                                                   InterPro;
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      136
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P19235; 1EBA.
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                                                                                                                                                                                                                                                                         SM00060;
                                                                                                        Similarity
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QRVEILEGRTECVLSNLRGRTRY 23
||||||:|||||:|||||| ||1
                                                                                                                                                                                                                                                    PS01352;
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                                                                                                                                                                                                                                                                                                               IPR003528;
                                                                                                                                                                                                                                                                                                                                   IPR008957;
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229 AA;
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229
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                                                                                  Conservative
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229
25196 MW;
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25170 MW;
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                                                                                                   88.9%;
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FN_III-like.
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01, Last sec
25, Last and
(Fragment).
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Last annotation updat
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Pred. No. 3.3e-09;
2; Mismatches 1
                                                                                  2;
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                                                                                  Score 104; DB Pred. No. 3.3e 2; Mismatches
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RESULT Q95N13

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Best Local S
Matches 20
                                                                                        P SEQUENCE FROM N.A.

A David B., Lim G.B., Moritz K.M., Koukoulas I., Wintour M.;

A David B., Lim G.B., Moritz K.M., Koukoulas I., Wintour M.;

T "Quantitation of the mRNA levels of Epo and EpoR in various tissue

I the ovine fetus.";

I Submitted (APR-2001) to the EMBL/GenBank/DDBJ databases.

R EMBL; AY029231; AAK38170.1; -.

R GO; GO:0016020; C:membrane; IEA.

R GO; GO:0016020; C:membrane; IEA.

R GO; GO:0004896; F:hematopoietin/interferon-class (D200-domain...

R GO; GO:0004872; F:receptor activity; IEA.

R InterPro; IPR003961; FN III.

R InterPro; IPR003962; FN III-like.

R InterPro; IPR003528; Hemitopoptn L-F1.

R Pfam, PF00041; fn3; 1.

R SMART; SM00060; FN3; 1.
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Q95N14;
01-DEC-2001
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Q95N13;
01-DEC-2001
01-DEC-2001
01-OCT-2003
Receptor.
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01-DEC-2001 (TrEMBLrel. 19, Last sequence up
01-OCT-2003 (TrEMBLrel. 25, Last annotation
Erythropoietin receptor (Fragment).
Ovis aries (Sheep)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ovis aries (Sheep).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Erythropoietin receptor (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the ovine fetus.";
Submitted (APR-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AN029232; AAK38737.1; -.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0004896; F:hematopoietin/interferon-class (D200-domain.
GO; GO:0004872; F:receptor activity; IEA.
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Mammalia; Eutheria; Ceta
Bovidae; Caprinae; Ovis.
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SMART; SM00060; FN3; 1
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InterPro; IPR008957; FN_III-like.
InterPro; IPR003528; Hemtopoptn_L_F1.
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                                                                          PS01352;
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387 AA;
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Pred. No. 5.9e-09;
2; Mismatches 1
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Best Local S
Matches 20
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STRAIN=AV19 / DSM 6324 / JCM 9639;

STRAIN=AV19 / DSM 6324 / JCM 9639;

K MEDLINE=21927647; PubMed=11930014;

K MEDLINE=21927647; PubMed=11930014;

K Slesarev A.I., Mezhevaya K.V., Makarova K.S., Polushin N.N.

A Shcherbinina O.V., Shakhova V.V., Belova G.I., Aravind L.,

A Natale D.A., Rogozin I.B., Tatusov R.L., Wolf Y.I., Stette

A Malykh A.G., Koonin E.V., Kozyavkin S.A.;

The complete genome of hyperthermophile Methanopyrus kand

T and monophyly of archaeal methanogens.";

Proc. Natl. Acad. Sci. U.S.A. 99:4644-4649(2002).

R EMBL, AE010332; AAM01560.1; -.

R InterPro; IPR009003; Cys_Ser_trypsin.
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Q8TYF5;
Q1-JUN-2002
01-JUN-2002
01-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                    Q9UQ52;
01-MAY-2000
01-MAY-2000
01-OCT-2003
EMBL, AB003592; BAA82612.1; -.
HSSP; P20241; 1CFB.
Genew; HGNC:2176; CNTN6.
GO; GO:0007155; P:cell adhesion; TAS.
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                                                                                                                             SEQUENCE FROM N.A.

MEDLINE-98146168; PubMed=9486763;
Kamei Y., Tsutsumi O., Taketani Y., Watanab
"cDNA cloning and chromosomal localization
                                                                                        molecule, NB-3 in human.";
J. Neurosci. Res. 51:275-283(1998)
                                                                                                                                                                                                                                                                    Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                          01.MAY-2000 (TrEMBLrel. 13, Created)
01.MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-CCT-2003 (TrEMBLrel. 25, Last annotation updat
Neural adhesion molecule NB-3.
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NCBI_TaxID=2320;
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                                                                                                                                                                                                                                                                    Chordata;
Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19285 MW; 81E181C7BF3BBD8B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 45282 MW;
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Last annotation update)
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Pred. No.
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Pred. No. 6.
                                                                                                                                                                                                                                                                    Craniata; Vertebrata; I Catarrhini; Hominidae;
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Q9AKX8;
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Pfam; PF00047; 19; 6.
SMART; SM00060; IGC2; 5.
SMART; SM004008; IGC2; 5.
PROSITE; PS00211; ABC TRANSPORTER 1; 1.
PROSITE; PS50835; IG_LIKE; 6.
                                                                                                                                       NB-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lueneberg E., Mayer B., Daryab N., Kooistra O., Zaehringer U., Rohde M., Swanson J., Frosch M.; "Chromosomal insertion and excision of a 30 kb instable genetic element is responsible for phase variation of lipopolysaccharide other virulence determinants in Legionella pneumophila."; Mol. Microbiol. 39:1259-1271(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR003439; ABC transporter.
InterPro; IPR003961; FN III.
InterPro; IPR008957; FN III-like.
InterPro; IPR007110; Ig-like.
InterPro; IPR003598; Ig_c2.
                                                                   Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hypothetical protein. SEQUENCE 826 AA; 91581 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteria; Proteobacteria; G
Legionellaceae; Legionella.
      SEQUENCE FROM N.A
                                        NCBI_TaxID=10116;
                                                                                                            Rattus norvegicus (Rat)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN=serogroup 1;
MEDLINE=21150471; PubMed=11251842;
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llarity 42.9%;
Conservative
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                                                                     Chordata;
Rodentia;
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Pred. No. 25;
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Pred. No. 9.9;
                                                                 Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Rat
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EMBL; D87248; BAA13320.1; -.
GG; GO:0016020; C:membrane; IEA.
GG; GO:0005524; F:ATP binding; IEA.
GG;:GO:0005524; F:ATP binding cassette (ABC) transporter acti. ..; IEA.
GG;:GO:000610; P:transport; IEA.
GO;:GO:000610; P:transport; IEA.
InterPro; IPR003499; ABC transporter.
InterPro; IPR003499; NBC III.
InterPro; IPR003499; NBC III-1ike.
InterPro; IPR007110; Ig-like.
InterPro; IPR007110; Ig-like.
InterPro; IPR007110; Ig-like.
InterPro; IPR007110; Ig-like.
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                                                                                                                                                                                                                  Submitted (OCT-2000) to the EMBL/GenBank/DDBJ databases. EMBL; AC011661; AAF16632.1; -. SEQUENCE 428 AA; 48982 MW; 603F3FF2725622F9 CRC64:
                                                                                                                                                                                                                                                                                                                                                          Shinn P., Brooks S., Buehler E., Chao Q., Johnson-Hopson C., Kh Kim C., Altafi H., Bei Q., Chin C., Chiou J., Choi E., Conn L., Conway A., Gonzales A., Hansen N., Howing B., Koo T., Lam B., Lienz C., Li J., Liu A., Liu K., Liu S., Mukharsky N., Nguyen M. Palm C., Pham P., Sakano H., Schwartz J., Southwick A., Thaveri Toriumi M., Vaysberg M., Yu G., Federspiel N.A., Theologis A., Ecker J.R.;
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Pfam; PF00047; ig; 6.

SMART; SM00060; FN3; 4.

SMART; SM00408; IGc2; 5.

PROSITE; PS00211; ABC TRANSPORTER_1; 1.

PROSITE; PS00815; IG_LIKE; 6.

Immunoglobulin domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Arabidopsis thaliana (Mouse-ear cress).

Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids;

eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                   "Genomic sequence for Arabidopsis thaliana BAC T23J18 from
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KRPQLLENTTLCILSITRGSRR
                                                   QRVEILEGRIECVLSNLRGRIR 22
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Pred. No.
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                                                                                                                                                              Score 48;
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., Thaveri A.,
                                                                                                                                                                                                                                                                                                                                      chromosome
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Matches 9
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Q8C6X1;
01-MAR-2003
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01-OCT-2003
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Q9JMB8;
01-OCT-2000
SEQUENCE FROM N.A.
STRAIN-C57BL/6J; TISSUB=Head;
MEDLINE=22354683; PubMed=12466851;
The FANTOM Consortium,
the RIKEN Genome Exploration Resea
                                                                                                                                                                                                                                                                                                                                                     SMART; SM00060; FN3; 4.
SMART; SM00408; IGC2; 5.
SMO0408; IGC2; 5.
PROSITE; PS00211, ABC_TRANSPORTER_1;
PROSITE; PS50835; IG_LIKE; 6.
PROSITE; PS50835; IG_MHC; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GO; GO:0016020; C:membrane; IEA.
GO; GO:0005524; F:ATP-binding cassette
GO; GO:0006810; P:transport; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Takahashi N., Watanabe K.; "Expression and regulation of a gene encoding molecule NB-3 of the contactin/F3 subgroup in Gene 245:253-266(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Neural recognition molecule CNTN6 OR MNB-3.
                                                                             Eukaryota; Metazoa;
Mammalia; Eutheria;
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Pfam; PF00047; ig; 6.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN=129SVJ; TISSUE=Brain;
MEDLINE=20183687; PubMed=10717476;
Lee S., Takeda Y., Kawano H., Hosoya H., Nomoto M., Fujimoto D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-OCT-2003
                                                               NCBI_TaxID=10090;
                                                                                                    Mus musculus
                                                                                                                 CNTN6.
                                                                                                                           Contactin
                                                                                                                                                                                                                                                                                                                                   SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro;
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                                                                                                                                                                                                                                                                                                                                              Immunoglobulin domain
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(TIEMBLrel. 15, Last sequence update)
(TIEMBLrel. 25, Last annotation update)
phition molecule NB-3.
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                                                                                                    (Mouse)
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 Exploration Research Group Phase I & II Team;
                                                                             Chordata;
Rodentia;
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Last sequence update)
Last annotation update)
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Pred. No. 47;
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                                                                            Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Mus
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Pfam; PF00047; ig; 6.

SMART; SM00060; FN3; 4.

SMART; SM00408; IG; 6.

SMART; SM00408; IG; 6.

PROSITE; PS00221; ABC TRANSPORTER 1; 1.

PROSITE; PS0035; IG LIKE; 6.

PROSITE; PS00290; IG MHC; 1.

SEQUENCE 1028 AA; 113761 MM; B233EI
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MGD; MGI:1858223; Cntn6.

GO:00016020; C:membrane; IEA.

GO; GO:0005524; F:ATP binding; IEA.

GO; GO:0005524; F:ATP-binding cassette

GO; GO:0006810; P:transport; IEA.

InterPro; IPR003439; ABC transporter.

InterPro; IPR00393; FN III.

InterPro; IPR003957; FN III.

InterPro; IPR003959; IN III-like.

InterPro; IPR003999; Ig.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Q94LS6;
01-DEC-2001
01-DEC-2001
                                                                                                                                                                                                                                       STRAIN=cv. Nipponbare;
Buell C.R., Yuan Q., Ouyang S., Moffat K.S., Hill J.N., Gansberg
Brenner M., Burgess S., Hance M., Shvartsbeyn M., Tsitrin T.,
Riggs F., Haiao J., Zismann V., Blunt S., Pai G., Vanaken S.E.,
Utterback T.R., Feldblyum T.V., Quackenbush J., Salzberg S.L.,
White O., Fraser C.M.;
"Oryza sativa chromosome 10 BAC OSJNBb0011A08 genomic sequence."
Submitted (MAY-2001) to the EMBL/GenBank/DDBJ databases.
EMBL, AC034258; AAK54287.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                              Hypothetical
SEQUENCE 52
                                                                                                                                                                                                                           Gramene; Q94LS6; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hypothetical protein. Oryza sativa (Rice).
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01-OCT-2003 (TrEMBLrel.
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Nature 420:563-573(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=4530;
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IPR003006;
                                                                                                                                                                          sal protein.
539 AA; Ko
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Ig-like.
Ig_c2.
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                                                                                       Score 47; DB Pred. No. 34; 2; Mismatches
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Pred. No.
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YL54 CAEEL
KKM1 RAT
FPG VIBPA
MURC CORGL
TSP3 MOUSE
TA29 TREPA
PTPD HUMAN
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HTII SALTY
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P32670 escherichia		P11141 caenorhabdi	P23759 homo sapien	P24610 mus musculu	P23760 homo sapien	P38867 saccharomyc	O29265 archaeoglob	Q9a2x6 caulobacter	060041 thermotoga	P77161 escherichia	

ALIGNMENTS

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15-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TISSUE=Placenta;

MEDLINE=92399733; PubMed=1668606;

Noguchi C.T., Bae K.S., Chin K., Wada Y., Schechter A.N.,

Hankins W.D.;

"Cloning of the human erythropoietin receptor gene.";

Blood 78:2548-2556(1991).
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Ehrenman K., St John T.;
"The erythropoietin receptor gene: cloning and identification
multiple transcripts in an erythroid cell line OCIMI.";
Exp. Hematol. 19:973-977(1991).
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                        PHOSPHORYLATION, AND INTERACTION WITH APS
MEDLINE=99301417; PubMed=10374881;
Wakioka T., Sabaki A., Mitsui K., Yokouch
                                                                                                                                                                                                                                                                                                                                                                                                  Chretien 8.; "Cloning of the gene encoding the human erythropoietin receptor."; Blood 78:2557-2563(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=92399734; PubMed=1668607;
Maouche L., Tournamille C., Hattab C., Boffa G., Cartron J.-P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=90304340; PubMed=2163696; Jones S.S., D'Andrea A.D., Halnes L.L., Williams exythropolectin receptor: cloning, characterization.";
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                                                               Yoshimura A.;
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   protein containing Pleckstrin homology (PH) and Src domains inhibits the JAK-STAT pathway in
                                                                                                                                                                                                                                                human erythropoietin
                                                                                           K., Yokouchi M.,
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                                                                                           Inoue A., Komiya S.,
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EMBL; M34986; AAA52401.1; EMBL; M60459; AAA52403.1; EMBL; M60459; AAA52393.1; EMBL; M76595; AAA52393.1; EMBL; M776595; AAA52392.1; EMBL; M77644; AAA52392.1; EMBL; M77644; AAA52392.1; EMBL; M77644; AAA52392.1; EMBL; AA3799; ZUHUR.

PDB; 1EBB; 29-UUL-97.
PDB; 1EBB; 29-UUL-99.
PDB; 1EBB; 10-CUT-99.
PDB; 1CM4; 11-AUG-99.
PDB; 1EBN; 07-JAN-00.
PDB; 1EBN; 07-JAN-00.
PDB; 1CN4; 11-AUG-99.
PDB; 1ENV; 07-JAN-00.
Genew; HGNC:3416; EPOR.
MIM; 133171; -.
GO; GO:0005887; C:integral to plasma membrane;
GO; GO:0004900; F:erythropoletin receptor activ
InterPro; IPR002996; CRIA.
InterPro; IPR008957; FN_III-like.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=98445092; PubMed=9774108; Syed R.S., Reid S.W., Li C., Cheetham J.C., Aoki K.H., Liu B., Zhan H., Osslund T.D., Chirino A.J., Zhang J., Finer-Moore J., Elliott S., Sitney K., Katz B.A., Matthews D.J., Wendoloski J.J., Eggrie J., Stroud R.M.; "Efficiency of signalling through cytokine receptors depends critically on receptor orientation.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Crystallographic evidence for preformed receptor before ligand activation."; Science 283.987-990(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             X-RAY CRYSTALLOGRAPHY (2.40 ANGSTROMS) OF 34-246.
MEDLINE=99141272; PubMed=9974392;
Livnah O., Stura E.A., Middleton S.A., Johnson D.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "An antagonist peptide-EPO receptor complex suggests dimerization is not sufficient for activation."; Nat. Struct. Biol. 5:993-1004(1998).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Livnah O., Stura E.A., Johnson D.L., Middleton S.A., Wrighton N.C., Dower W.J., Jolliffe L.K., Wilson I.A "Functional mimicry of a protein hormone by a peptid
                                                                                                                                                                                                                                                                                                                                  entities requires a license agreement (So or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EPO
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Leukemia 13:760-767(1999).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FUNCTION: Receptor for erythropoietin. Mediates erythropoietin-
induced erythroblast proliferation and differentiation. Upon EPO
stimulation, EPOR dimerizes triggering the JAK2/STAT5 signaling
cascade. Isoform 2 acts as a dominant-negative receptor of EPOR-
mediated signals.
SUBUNIT: Interacts with APS.
SUBCELLULAR LOCATION: Type I membrane protein.
SIMILARITY: Belongs to the type I cytokine family of receptors.
SUBULLARITY: Contains 1 fibronectin type III domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               receptor complex at 2.8 A.";
ence 273:464-471(1996).
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W., Krause C.D.,
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PIRSF; PIRSF001959; EPO_receptor;
SMART; SM00060; FN3; 1.
             Eukaryota; Metazoa;
Mammalia; Eutheria;
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mbrane; Glycoprotein;
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Rodentia;
                                                                                                                                                                                                                                                                                                                                          55065 MW;
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                                                                                                                                                                                                                                                                                            Score 117;
Pred. No. 8.
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N-LINKED (GLCNAC.
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             Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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                Mus
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D'Andrea A.D., Lodish H.F., "Expression cloning of the Cell 57:277-285 (1989).
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or send a
EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Spleen focus-forming virus long terminal activation of the murine erythropoietin rufriend leukemia cell line.";
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MEDLINE=92017832; PubMed=1656233;

Hino M., Tojo A., Misawa Y., Morii H., Takaku

"Unregulated expression of the erythropoietin

insertion of spleen focus-forming virus long t

murine erythroleukemia cell line.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "The adapter protein APS associates with the multifunctional docking sites Tyr-568 and Tyr-936 in c-Kit."; Biochem. J. 370:1033-1038(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gisselbrecht S., Cartron J.-P
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"Characterization of murine erythropoietin
J. Mol. Biol. 216:567-575(1990).
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MEDLINE=91080149; PubMed=2175360;
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SEQUENCE FROM N.A.
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                                                                                           s SWISS-PROT entry is copyright. It is produced through a collaboration ween the Swiss Institute of Bioinformatics and the EMBL outstation - Buropean Bioinformatics Institute. There are no restrictions on its by non-profit institutions as long as its content is in no way ified and this statement is not removed. Usage by and for commercial ities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                    FUNCTION: Receptor for erythropoietin. Mediates erythropoietin-
induced erythroblast proliferation and differentiation. Upon EPO
stimulation, EPOR dimerizes triggering the JAK2/STAT5 signaling
cascade. Isoform 2 acts as a dominant-negative receptor of EPOR-
mediated signals.
SUBCELLULAR ICCATION: Type I membrane protein.
SUBCELLULAR ICCATION: Type I membrane protein.
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  J04843;
X53081;
                                                                      an email to license@isb-sib.ch).
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  AAA37571.1;
CAA37248.1;
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Zon L.I., Orkin S.H.,
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f the murine erythropoietin receptor.";
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InterPro; IPR008957; FN_III-like.
InterPro; IPR003961; FN_III.
InterPro; IPR003961; Hemtopoptn_L_F1.
Pfam; PF00041; fn3; 1.
PIRSF; PIRSF001959; EPO_receptor; 1.
SMART; SM00060; FN3; 1.
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EMBL; M62360; AAA3758;
EMBL; S59388; AAB2002:
PIR; A41686; A32385.
PIR; S14081; S14081.
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
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                                                                                                                                                                                                                J. Biol. Chem. -i- FUNCTION: 1
                                                                                                                                                                                                                                                      cells."
                                                                                                                                                                                                                                                                      "Functional erythropoietin racharacteristics. Comparison
                                                                                                                                                                                                                                                                                                          Masuda S., Nagao M.,
Tabira T., Sasaki R.;
                                                                                                                                                                                                                                                                                                                              MEDLINE=93266574; PubMed=7684373; Masuda S., Nagao M., Takahata K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Erythropoietin receptor precursor
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MGD; MGI:95408; Epor.
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                                                                           cascade. Isoform 2 acts
mediated signals.
- SUBCELLULAR LOCATION: T
- SIMILARITY: Belongs to
Subfamily 1.
                                                             SIMILARITY: Contains 1 fibronectin type III domain.
                                                                                                                                                         FUNCTION: Receptor for erythropoietin. Mediates erythropoietin-
induced erythroblast proliferation and differentiation. Upon EPG stimulation, EPOR dimerizes triggering the JAKZ/STATS signaling cascade. Isoform 2 acts as a dominant negative receptor of EPOR.
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Q8G674;
10-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Receptor;
This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                        10-OCT-2003 (Rel. 42, 0
10-OCT-2003 (Rel. 42, 1
10-OCT-2003 (Rel. 42, 1
Methyltransferase gidB
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TRANSMEM
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                                                                                                                                                                                                                                                                                                                                                                                                   BIFLO
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                                                                                                                                                               Pridmore R.D., Arigoni
                                                                                                                                                                     MEDLINE=22294977; PubMed=12381787;
Schell M.A., Karmirantzou M., Snel B., Vilanova D., Berg
Pessi G., Zwahlen M.-C., Desiere F., Bork P., Delley M.,
Pridmore R.D., Arigoni F.;
                                                                                                                                                                                                                                                                           Bifidobacterium longum.
Bacteria, Actinobacteria,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
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                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                   Bifidobacteriaceae; Bifidobacterium
                                                                                                                                                                                                                                                                                                    GIDB OR BL0646.
                                                                                                                                                                                                                                                                                                                 protein B)
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                                                                                                                                                                                                                                                    NCBI_TaxID=216816;
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                                                                                                                                               the
                                                                                                       e genome sequence of Bifidobacterium longum refl
the human gastrointestinal tract.";
c. Natl. Acad. Sci. U.S.A. 99:14422-14427(2002).
FUNCTION: probable S-adenosyl-L-methionine depen
methyltransferase specific for a sterol and/or
                                                                                  SIMILARITY: Belongs to the gidB family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A46713; A46713.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P19235; 1EBA.
Pro; IPR002996; CR1A.
Pro; IPR008957; FN III-like.
Pro; IPR003966; FN III.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SM00060; FN3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PIRSF001959; EPO_receptor; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 QRVEVLEGRIECVLSNLRGGTRY 215
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Transmembrane;
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Last
B (EC
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BY SIMILA
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                                                                                                                                                                                                                                                                              Actinobacteridae; Bifidobacteriales;
                                                                                                                                                                                                                                                                                                                           sequence update)
annotation update)
2.1.-.-) (Glucose
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ERYTHROPOIETIN RECEPTOR.
EXTRACELLULAR (POTENTIAL).
POTENTIAL.
CYTOPLASMIC (POTENTIAL).
FIBRONECTIN TYPE-III.
BY SIMILARITY.
BY SIMILARITY.
BY SIMILARITY.

N-LINKED (GLCNAC. . .) (POTE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred. No. 1.5
L; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 109;
Pred. No. 1
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AC79AF22D06A7312 CRC64;
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.5e-09;
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                                                                                                         ine dependent and/or lipid
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                                                                                                                                                          reflects
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                                                                                                                                                          its adaptation
                                                                                                           substrate
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DEPENDENCE OF THE PROPERTY OF

InterPro; [nterPro;

GO:0016021; C:int erPro; IPR000859;

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RESULT 5
KRM1_MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                       "Molecular cloning and characterization of Kremen, a novel

kringle-containing transmembrane protein.";

Biochim. Biophys. Acta 1518:63-72(2001).

c |- FUNCTION: Receptor for Dickkopf protein. Cooperates with Dickkopf

to block Wnt/beta-catenin signaling (By similarity).

c |- SUBCELLULAR LOCATION: Type I membrane protein (Potential).

c |- TISSUE SPECIFICITY: In the adult, widely expressed with high

c |- Levels in heart, lung, kidney, skeletal muscle and testis.

c |- DEVELOPMENTAL STAGE: In the embryo, expression is first detected

c |- DEVELOPMENTAL STAGE: In the embryo, expression is first detected

c |- adult, At 9.5 dpc, expression is localised to the apical

c |- adult, At 9.5 dpc, expression is localised to the apical

c |- etcodermal ridge (AER) of the developing fore- and hindlimb buds,

the telencephalon and the first brachial arch. At 10.5 dpc,

expression is also observed in the myotome and in sensory tissues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KRM1 MOUSE
Q99N43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUE=Brain, and Kidney;
MEDLINE=21167372; PubMed=11267660;
Nakamura T., Aoki S., Kitajima K.,
                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Entropean Bioinformatics Institute. There are no restue by non-profit institutions as long as its content use by non-profit institutions as long. Its content use by non-profit institutions as long as its content use by non-profit institutions as long.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TIGREAMS; TIGRO0138; gidB; 1.
Transferase; Methyltransferase;
SEQUENCE 221 AA; 24163 MW; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; AE014686; AAN24468.1; -. HAMAP; MF_00074; -; 1. Interpro; IPR003682; GidB. Pfam; PF02527; GidB; 1.
                                                                       EMBL; AB059617;
HSSP; P00747; 10
                                                                                                                                               entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and the nose) (Dickkopf receptor).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kremen protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-FEB-2003
10-OCT-2003
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                                           MGB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nakamura T.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.,
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                                                                                                                                                                                                                                                                                                                                              such as the nasal pit and optic vesicle.
SIMILARITY: Contains 1 CUB domain.
SIMILARITY: Contains 1 kringle domain.
SIMILARITY: Contains 1 WSC domain.
MGI:1933988; Kremen.
GO:0016021; C:integral to
rPro; IPR000859; CUB.
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(Rel. 42, Last annotation update)
in 1 precursor (Kringle-containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                          1CEA.
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                                                                                                 BAB40968.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chordata;
Rodentia;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20
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MBL outstation -
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Best Local S
Matches 11
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PRODOM; PD000395; Kringle; 1.
SMART; SM00042; CUB; 1.
SMART; SM00130; KR; 1.
PROSITE; PS01180; CUB; 1.
PROSITE; PS00021; KRINGLE 1; 1.
PROSITE; PS00021; KRINGLE 2; 1.
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SEQUENCE
                                                                                                                                                                                            SEQUENCE FROM N.A. (ISOFORM 2).

Tashiro H., Yamazaki M., Watanabe K., Kumagai A., Itakura S.,

Pukuzumi Y., Fujimori Y., Komiyama M., Sugiyama T., Irie R.,

Otsuki T., Sato H., Wakamatsu A., Ishii S., Yamamoto J., Isono Y.,

Kawai-Hio Y., Saito K., Nishikawa T., Kimura K., Yamashita H.,

Katana K., Nakamura Y., Sekine M., Kikuchi H., Kanda K., Wagatsuma M.,

Murakawa K., Kanehori K., Takahashi-Fujii A., Oshima A., Sugiyama A.,

Kawakami B., Suzuki Y., Sugano S., Nagahari K., Masuho Y., Nagai K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      KRM1 HUMAN STANDARD; PRT; 475 AA. Q96MUB; Q9BY70; Q9UGS5; Q9UGU1; 28-FEB-2003 (Rel. 41, Created) 28-FEB-2003 (Rel. 41, Last sequence update) 10-OCT-2003 (Rel. 42, Last annotation update) Kremen protein 1 precursor (Kringle-containing and the nose) (Dickkopf receptor).
KREMENI OR KREMEN.
        MEDLINE=20057165; PubMed=10591208; Dunham I., Hunt A.R., Collins J.E., Clamp M., Smink L.J., Ainscough R., Bagguley C., Bailey J., Barlow K.F.,
                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A. (I)
Nakamura T., Nakamura
Submitted (OCT-2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CHAIN
DOMAIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DOMAIN
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                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DOMAIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11;
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                                                                                                                                         (OCT-2001) to the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (Human)
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                                                                                                                                                                                                                                                                                                                                                                                                (ISOFORM nura T.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                41.9%;
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N-LINKED (GLCNAC...

N-LINKED (GLCNAC...
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                                                                                                                                      project.";
EMBL/GenBank/DDBJ
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Pred. No.
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CYTOPLASMIC (POTENTIAL)
KRINGLE.
WSC.
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EXTRACELLULAR (POTENTIAL)
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Bruskiewich R., Beart
Almeida J.P., Babbage
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                             R., Beare D.M., ., Babbage A.K.,
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RA Burrill W.D. Burton J. Carder C. Carter N.E. Commor R.
RA Clegg S.M., Cobley V.E., Cole C.G., Collier R.E., Connor R.
RA Clay D., Corby N.R., Coville G.J., Cox A.V., Davis J., Dawson E.,
RA Conroy D., Corby N.R., Coville G.J., Cox A.V., Davis J., Dawson E.,
RA Brans K.L., Fey J.M., Fleming K., French L., Garner A.A.,
RA Evans K.L., Fey J.M., Fleming K., French L., Garner A.A.,
RA Hall R.E., Hall-Tamlyn G., Heathcott R.W., Ho S., Holmes S.,
RA Hall R.E., Jones M.C., Kershaw J., Kimberley A.M., King A.,
RA Hart S.E., Jones M.C., Kershaw J., Kimberley A.M., King A.,
RA Martyn I.D., Mashreghi-Mohammadi M., Matthews L.H., Mccann O.T.,
RA Martyn I.D., Mashreghi-Mohammadi M., Matthews L.H., Mccann O.T.,
RA Mclay J., Mclaren S., McMurray A.A., Milne S.A., Mortimore B.J.,
RA Mclay J., Mclaren S., Kenchaw J., Kimberley A.M., Mortimore B.J.,
RA Mclay J., Mclaren S., Kuce C.D., Smalley S., Smith M.L.,
RA Mclay J., Melaren S., Kuce C.D., Smalley S., Smith M.L.,
RA Walliams L., Williams S.A., Williamson H., Wilmer T.E., Wilming L.,
RA Williams L., Williams S.A., Williamson H., Mitsuyama S.,
RA Williams S., Kawasaki K., Sasaki T., Asakawa S., Kudoh J.,
RA Williams J., Shibuya K., Yoshizaki Y., Aoki N., Mitsuyama S.,
RA Minoshima S., Kawasaki K., Sasaki T., Asakawa S., Kudoh J.,
RA Williams J., Levis S., Lin S.-P., Loh P., Malaj E., Nguyen T., Pan H.,
RA Mand A., Fang F., Fu Y., Hu P., Hua A., Kenton S., Lai H., Lao H.I.
RA Mand N., Johnson D., Bemis G., Bentley D., Bradshaw H., Bourne S.,
RA Mann M., Zhang G., Chissoe S., Murray J., Miller N., Mins P.,
RA Mann M., Zhang G., Chissoe S., Murray J., Miller N., Mins P.,
RA Mann M., Johnson D., Bemis G., Bentley D., Bradshaw H., Bourne S.,
RA Minoshima S., Kambar Y., Wang Y., W
                                                                                                                                                                                           EMBL;
                                                                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                        or send
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <del>-</del>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nature 402:489-495(1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                        NOTE=NO EXI-
                                                                                                                                                                                                                                                                                                                                                                                                                             SIMILARITY: Contains 1 CUB domain. SIMILARITY: Contains 1 kringle domain. SIMILARITY: Contains 1 WSC domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FUNCTION: Receptor for Dickkopf protein. Cooperates with Dickkopf to block Wnt/beta-catenin signaling (By similarity).
SUBCELLULAR LOCATION: Type I membrane protein (Potential).
ALTERNATIVE PRODUCTS:
Event=Alternative splicing; Named isoforms=2;
                                                                                                                                                                                        AB059618; BAB40969.1; -. AK056425; BAB71180.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IsoId=Q96MU8-1; Sequence=Displayed;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IsoId=Q96MU8-2; Sequence=VSP_003900
                                                                                                                                                                                                                                                           an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Blakey S.E., Bridgeman A.M., Buck
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   experimental confirmation available;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Carter N.P., Chen Y.,
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EMBL; AL021393; CAB62959.1; -. Genew; HGNC:17550; KREMENI. GO; GO:0016021; C:integral to membrane; GO; GO:0005524; C:membrane fraction; TAS GO; GO:0007154; P:cell communication; TA

Z95116; CAB62952.1;

InterPro; (nterPro;

; IPR000859; ; IPR000001; ; IPR002889;

Kringle. WSC. CUB.

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SOUTH THE TENT TO 
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Best Local (
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10-OCT-2003
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CONFLICT
SEQUENCE
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CARBOHYD
CARBOHYD
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DOMAIN
TRANSMEM
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CARBOHYD
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CARBOHYD
                                                                                                                                                                                Nishio Y., Nakamura Y., Kawarabayasi Y., Usuda Y. Sugimoto S., Matsui K., Yamagishi A., Kikuchi H., Gojobori T.;
                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
STRAIN=YS-314 / AJ 12310 / DSM 44549 / JCM 11189;
MEDLINE=22723752; PubMed=12840036;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COREF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PROSITE; PS01180; CUB; 1.
PROSITE; PS00121; KRINGLE_2; 1.
PROSITE; PS50070; KRINGLE_2; 1.
Wht signaling pathway; Signal;
Alternative splicing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRINTS; PRO0018; KRINGLE.
PRODOMS, PD000395; Kringle; 1.
SMART; SM00042; CUB; 1.
SMART; SM00130; KR; 1.
                                                                                                                          replacements responsible efficiens.";
                                                                                                                                                                                                                                                                                                                                Bacteria; Actinobacteria; Actinobacteri
Corynebacterineae; Corynebacteriaceae;
                                                                                                                                                                                                                                                                                                                                                                                        transcarbamylase) (ATCase).
PYRB OR CE1732.
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                                                                                                                                                                                                                                                                                                             NCBI_TaxID=152794;
                                                                                                                                                                                                                                                                                                                                                                      Corynebacterium efficiens.
                                                                                                                                                                                                                                                                                                                                                                                                                            Aspartate carbamoyltransferase (EC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DOMAIN
                                                                                                                                                                 "Comparative complete
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7
                                PATHWAY: Pyrimidine biosynthesis; second step. SIMILARITY: Belongs to the ATCase/OTCase family.
                                                                      CATALYTIC ACTIVITY: Carbamoyl phosphate + L-aspartate + N-carbamoyl-L-aspartate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PF00431;
PF00051;
PF01822;
SWISS-PROT entry
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                475 AA;
                                                                                                           13:1572-1579(2003).
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206
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395
416
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121
121
216
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219
235
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347
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475
394
415
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116
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323
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 18
                                                                                                                                             genome sequence analysis of the amino acid ble for the thermostability of Corynebacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Created)
copyright.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MW;
                                                                                                                                                                                                                                                                                                                                  Actinobacteridae; Actinomycetales; bacteriaceae; Corynebacterium.
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KREMEN PROTEIN :

EXTRACELLULAR ()
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 49;
Pred. No.
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I -> V (IN REF. 2).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CYTOPLASMIC (POTENTIAL)
KRINGLE.
WSC.
CUB.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  B7E86FD80F96A0A4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                          ion update)
2.1.3.2) (Aspartate
 It is
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4.1;
                                                                                                                                                                                                                    Usuda Y.,
produced through
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RESULT 8
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                                                  HAMAP; MF_00001; -; 1.
InterPro; IPR006130; As
InterPro; IPR0082; As
InterPro; IPR06131; OT
InterPro; IPR066131; OT
InterPro; IPR066132; OT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-FEB-2003
28-FEB-2003
10-OCT-2003
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InterPro; IPR006130; Asp/Orn_COtranf.
InterPro; IPR002032; Asp_carbmltransf.
InterPro; IPR006131; OTCace_O.
InterPro; IPR006132; OTCace_P.
Pfam; PF00185; OTCace; 1.
Pfam; PF02729; OTCace_N; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the European Bloinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EW the EWropean Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-s
                                                                                                                                                                                                                                                                                                              "Complete genomic sequence of Corynebacterium glutamicum ATC Submitted (MAY-2002) to the EMBL/GenBank/DDBJ databases.
-i- CATALYTIC ACTIVITY: Carbamoyl phosphate + L-aspartate =
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TIGREAMS; TIGRO0670; asp_carb_tr; 1.
PROSITE; PS00097; CARBAMOYLTRANSFERASE; 1.
Pyrimidine biosynthesis; Transferase; Comp
             Pfam; PF00185; OTCace; 1.
Pfam; PF02729; OTCace N; 1.
PRINTS; PR00100; AOTCASE.
                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
STRAIN=ATCC 13032 /
                                                                                                                                                                                                                                                                                                                                                                                                                                             Corynebacterium glutamicum (Brevibacterium flavum).
Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Aspartate carbamoyltransferase transcarbamylase) (ATCase).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            QBNQ38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PYRB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
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                                                                                                                          EMBL; AP005279; BAB99005.1; -.
                                                                                                                                                       or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                           Nakagawa S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                Corynebacterineae; Corynebacteriaceae; Corynebacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PYRB OR CGL1612.
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                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=1718;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                       PATHWAY: Pyrimidine biosynthesis; second step. SIMILARITY: Belongs to the ATCase/OTCase family.
                                                                                                                                                                                                                                                                                                   + N-carbamoyl-L-aspartate.
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(Rel. 41, Last sequence update)
(Rel. 42, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               33706 MW;
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                                                                    Asp/Orn_COtranf.
Asp_carbmltransf.
OTCace_O.
                                                     OTCace_P
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2.1.3.2) (/
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(See http://www.isb-sib.
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databases.
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asp_carb_tr; 1.

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TSP3_HUMAN STANDARD; PRT; 956
AC P49746, OBWV34,
DT 01-CCT-1996 (Rel. 34, Last sequence updat
DT 01-CCT-1996 (Rel. 34, Last sequence updat
DT 01-CCT-2003 (Rel. 42, Last annotation upo
DE DT 01-CCT-2003 (Rel. 42, Last annotation upo
DE Thrombospondin 3 precursor.

OS HOMO sapiens (Human).

OC ENLARYOTE, Metazoa; Chordata; Craniata; '
OC Mammalia; Eutheria; Primates; Catarrhini
OX NCBI_TAXID=9606;
RN [1]
RN SEQUENCE FROM N.A.
RC TISSUB-Lung;
RX MEDLINE=96044440; PubMed=7558000;
RA Adolph K.W., Long G.L., Winfield S., Gin
RT "Structure and organization of the human
RT Genomics 27:329-336(1995).
RN [2]
RN [2]
RN SEQUENCE OF 1-736 FROM N.A.
RX MEDLINE=97474796; PubMed=9331372;
RA MEDLINE=97474796; PubMed=9331372;
RA MIDIANE=STANDARD; PubMed=933172;
RA MIDIANE=STANDARD; PubMed=933172;
RA MIDIANE=STANDARD; PubMed=933172;
RA MEDLINE=97474796; PubMed=933172;
RA MIDIANE=STANDARD; PubMed=933172;
RA MEDLINE=97474796; PubMed=12477932;
RA SEQUENCE OF 365-956 FROM N.A.
RX MEDLINE=22388257; PubMed=12477932;
RA SEQUENCE OF 365-956 FROM N.A.
RC TISSUB-Retina;
RA SEQUENCE OF 365-956 FROM N.A.
RC TISSUB-Retina;
RA Diatchenko L., Marusina K., Farmer A.A.
RA SEQUENCE OF 365-956 FROM N.A.
RA SLEECHUL S.F., Zeeberg B., Buetow K.H., SA
RA Altechul S.F., Jordan H., Moore T., MAA
RA SLEECHUL S.F., Jordan H., Moore T., MAA
RA SLEECHUL S.F., Jordan H., Moore T., MAA
RA BLOWNSEEN M.J., WGEWAN P.J., WEKERAN K.J., RA
BOORAS S.A, McEwan P.J., WEKERAN K.J., Shevch
RA BLOWNSEEN M.J., WGEWAN J.W., Sodergren E.J.
RA RICHARDS S., WOTLEY K.C., Hale S., Garcii
RA Whiting M., MAAAN A., Young A.C., Shevch
RA BLOWNSEEN A., Schein J.E., Jones S.J.M., if
RI Fahay J., Helton E., Kettenan M., Madan, J.
RA RICHARDS S., WOTLEY K.C., Hale S., Garcii
RA Whiting M., MAAAN A., Young A.C., Shevch
RA BLOWNSEEN A., Schein J.E., Jones S.J.M., if
RA BLOWNSEEN A., Schein J.E., Jones S.J.M., if
RA BLOWNSEEN A., Schein J.E., Jones S.J.M., if
RA Hopkins R.G., Shevch
RA BLOWNSEEN A., Schein J.E., Jones S.J.M., if
RA GARCING S., WOTLEY K.C., Hale S., Garcii
RA Hopkins R.G.,
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RA Strausberg R.D., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Alteschul S.F., Zeeberg B., Buetcw K.H., Schaefer C.F., Bhat N.K.,
RA Alteschul S.F., Zeeberg B., Buetcw K.H., Schaefer C.F., Bhat N.K.,
RA Alteschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Biatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Biatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahby J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwoo
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Best Local
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4 MW; F33154F4D033CF22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   artin B.M., Ginns E.I., Sidransky E.; ditional genes contiguous to the chromosome lq21: implications for Gaucher
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           field S., Ginns E.I., Bornsteir of the human thrombospondin 3
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Pred. No. 3
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3.7;
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           GG; GO:0005509; F:calcium ion binding; TAS
GO; GO:0007160; P:calcium ion binding; TAS
GO; GO:0007160; P:call-matrix adhesion; TA
InterPro; IPR006209; EGF_Ca.
InterPro; IPR006209; EGF_Like.
InterPro; IPR003367; tsp_3.
InterPro; IPR003367; tsp_3.
InterPro; IPR003129; TSPG.
InterPro; IPR003129; TSPN.
Pfam; PF00008; EGF; 1.
Pfam; PF00008; EGF; 1.
Pfam; PF007315; TSPG; 1.
Pfam; PF02710; TSPN; 1.
SMART; SM00179; EGF CA; 1.
SMART; SM00179; EGF CA; 1.
PROSITE; PS01186; EGF 2; 1.
PROSITE; PS01186; EGF 2; 1.
PROSITE; PS01186; EGF CA; 2.
Glycoprotein; Cell adhesion; Calcium-bindi
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DOMAIN
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MIM; 188062; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; L38969; AAC41762.1; -.
EMBL; AF023368; AAC51818.1; -
EMBL; BC018786; AAH18786.1; -
EMBL; BC18786; A57121.
PIR; A57121; A57121.
HSSP; P35555; IENN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entities requires a or send an email to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SIGNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tities requires a license agreement (S send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GO: 0005509;
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           956
272
315
369
413
       EGF-LIKE 1.
EGF-LIKE 2, CAI
EGF-LIKE 3, CAI
EGF-LIKE 4.
TSP TYPE-3 1.
TSP TYPE-3 2.
TSP TYPE-3 3.
TSP TYPE-3 5.
TSP TYPE-3 5.
TSP TYPE-3 6.
TSP TYPE-3 7.
C-TERMINAL.
INTERCHAIN (PRO
INTERCHAIN (PRO
BY SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Calcium-binding;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 binding; TAS. adhesion; TAS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (PROBABLE)
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(POTENTIAL)

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Paulsen I.T., Banerjei L., Myers G.S.A., Nelson K.E., Seshadri R.,
Paulsen I.T., Banerjei L., Myers G.S.A., Heidelberg J.F.,
Read T.D., Fouts D.E., Eisen J.A., Gill S.R., Heidelberg J.F.,
Tettelin H., Dodson R.J., Umayam L., Brinkac L., Beanan M.,
Daugherty S., DeBoy R.T., Durkin S., Kolonay J., Madupu R., Nelson W
Vamathevan J., Tran B., Upton J., Hansen T., Shetty J., Khouri H.,
Utterback T., Radune D., Ketchum K.A., Dougherty B.A., Fraser C.M.;
"Role of mobile DNA in the evolution of vancomycin-resistant
Enterococcus faecalis.";
Science 299:2071-2074(2003).

-i- SIMILARITY: Belongs to the tautomerase family.
                                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                   Isomerase; Complete proteome.
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HAMAP; MF_00718; -; 1.
InterPro; IPR004370; T
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STRAIN=V583 / ATCC 700802;
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Bacteria; Firmicutes; Lactobacillales; Enterococcaceae; Enterococcus.
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RESULT 11 RS28_KLULA

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RESULT 12
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P33285;
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InterPro; IPR000289; Ribosomal_
Pfam; PF01200; Ribosomal_S28e;
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Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Kluyveromyces.
           This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its
                                                                               Hoekstra R., Ferreira P.M., Bootsman T.C., Mager W.H., Pla "Structure and expression of the ABF1-regulated ribosomal gene in Kluyveromyces."; Yeast 8:949-959(1992).
                                                                                                                                        STRAIN=CBS 6556; MEDLINE=93127729; PubMed=1481571;
                                                                                                                                                                                            Kluyveromyces marxianus (Yeast) (Kluyve
Eukaryota; Pungi; Ascomycota; Saccharon
Saccharomycetales; Saccharomycetaceae;
NCBI_TaxID=4911;
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ProDom; PD005541; Ribosomal_S28e; 1.
PROSITE; PS00961; RIBOSOMAL_S28E; 1.
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Yeast 8:949-959(1992).
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    SIMILARITY: Belongs to the S28E family of ribosomal proteins.

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al protein S28 (S33).
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Q59572;
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MEDLINB=22206494; PubMed=12218036;

Fleischmann R.D., Alland D., Eisen J.A., Carpenter L., White O.,

Peterson J., DeBoy R., Dodson R., Gwinn M., Haft D., Hickey E.,

Kolonay J.F., Nelson W.C., Umayam L.A., Ermolaeva M., Salzberg S.L.,

Kolonay J.F., Nelson W.C., Umayam L.A., Ermolaeva M., Salzberg S.L.,

Bolcher A., Utterback T., Weidman J., Khouri H., Gill J., Mikula A.,

Bishai W., Jacobs W.R. Jr., Venter J.C., Fraser C.M.;

"Whole-genome comparison of Mycobacterium tuberculosis clinical and
laboratory strains.";
                                                                                                                                                                                                                                                                                                                                                                                                                                    SPECIES-M.tuberculosis; STRAIN-H37Rv;

REDIINE-98295987; PubMed=9634230;

Cole S.T., Brosch R., Parkhill J., Garnier T., Churcher C., Harris Gordon S.V., Eiglmeier K., Gas S., Barry C.E. III, Tekaia F., Gordon S.V., Eiglmeier K., Gas S., Barry C.E. III, Tekaia F., Badcock K., Basham D., Brown D., Chillingworth T., Connor R., Davies R., Devlin K., Feltwell T., Gentles S., Hamlin N., Holroyd Hornsby T., Jagels K., Krogh A., McLean J., Moule S., Murphy L., Oliver S., Osborne J., Quail M.A., Rajandream M.A., Rogers J., Rutter S., Seeger K., Skelton S., Squares S., Squares R., Sulston J.E., Taylor K., Whitehead S., Barrell B G.;

"Deciphering the biology of Mycobacterium tuberculosis from the complete genome sequence.";
Garnier T., Eiglmeier K., Camus J.-C., Medina N., Mansoor H., Pryor M., Duthoy S., Grondin S., Lacroix C., Monsempe C., Simon S. Harris B., Atkin R., Doggett J., Mayes R., Keating., Wheeler P.I. Parkhill J., Barrell B.G., Cole S.T., Gordon S.V., Hewinson R.G., "The complete genome sequence of Mycobacterium bovis."; Proc. Natl. Acad. Sci. U.S.A. 100:7877-7882 (2003).
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01-NOV-1997 (Rel. 35, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Putative cytochrome P450 128 (EC 1.14.-.-).
CYP128 OR RV2268C OR MT2330 OR MTCY339.42 OR MB2291C.
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PIR; S30006; S30006.
InterPro; IPR000289; Ribosomal S28e.
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                                                                                                                SPECIES=M.bovis; STRAIN=AF2122/97; MEDLINE=22709107; PubMed=12788972;
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Monsempe C., Simon S.,
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RESULT 14
AMS2_SCHPO
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AX MEDILINE-21849401; PubMed=11859360;

REDILINE-21849401; PubMed=11859360;

RA Sgouros J., Feat N., Hayles J., Baker S., Basham D., Bowman S., Sarokas K., Brown D., Brown S., Chillingworth T., Churcher C.M., Commor R., Cronin A., Davis P., Feltwell T., Fraser A., Gentles S., Goble A., Hamlin N., Harris D., Hidalgo J., Hodgson G., A., Holroyd S., Hornsby T., Howarth S., Huckle E.J., Hunt S., Jagels K., A., Holroyd S., Hornsby T., Howarth S., McDonald S., McLean J., R., Money P., Moule S., Mungall K., Murphy L., Niblett D., Odell C., R., Money P., Moule S., Mungall K., Murphy L., Niblett D., Odell C., R., A., O'Neil S., Pearson D., Quail M.A., Rabbinowitsch E., Ra, Multerford K., Rutter S., Saunders D., Seeger K., Sharp S., R., Skelton J., Simmonds M., Squares R., Squares S., Stevens K., Taylor K., Taylor R.G., Tivey A., Walsh S.V., Warren T., Whitehead S., R., Woodward J., Volckaert G., Aert R., Robben J., Grymonprez B., Weltjens I., Vanstreels E., Rieger M., Schaefer M., Mueller-Auer S., Ra, Gabel C., Fuchs M., Fritzc C., Holzer E., Moestl D., Hilbert H.,
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                           10-OCT-2003 (Rel. 42, Created)
10-OCT-2003 (Rel. 42, Last sequence up
10-OCT-2003 (Rel. 42, Last annotation
CENP-A multicopy suppressor protein 2
AMS2 OR SPCC290.04 OR SPCC4F11.01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; BX248342; PIR; H70729; H70729.
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Q9URT4; Q9P7Z1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entities requires a license agreement (S or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -!- SUBCELLULAR LOCATION: Integral membrane protein.
                                                                                                                                                                                                                                                                                                                                                              Schizosaccharomyces pombe (Fission yeast).
Eukaryota; Fungi; Ascomycota; Schizosaccharomycetes;
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                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
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MT2330; -.
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Pred. No.
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POTENTIAL.

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IRON (HEME AXIAL LIGAND) (BY SIMILARITY).

633F233CEF03AD7A CRC64;
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Matches
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EMBL; AL117389; CAB55767.2; -.
GenebB_SPombe; SPCC290.04; -.
GO; GO:0005694; C:chromosome; ISS.
GO; GO:0005654; C:nucleoplasm; ISS.
GO; GO:0007049; P:cell cycle; ISS.
InterPro; IPR000679; Znf_GATA.
                                                                                                                                                                                                                                                                                                -YL54 CAEEL
P34434;
01-FEB-1994
01-FEB-1994
28-FEB-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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Chen E.S., Saltoh S., Yanagida M., Takahashi K.;
"A cell cycle-regulated GATA factor promotes centromeric localization of CENP-A in fission yeast.";
Mol. Cell 11:175-187(2003).
STRAIN=Bristol N2;
MEDLINE=94150718; PubMed=7906398;
                                                                                                                                                   Eukaryota; Metazoa; Nemat
Rhabditidae; Peloderinae;
                                                                                                                                                                                                                                                              01-FEB-1994 (Rel. 28, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Hypothetical protein F44E2.4 in chromosome III.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   CAEEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PROSITE; PS00344; GATA ZN FINGER 1; FALSE NEG. PROSITE; PS50114; GATA ZN FINGER 2; 1. Transcription regulation; Nuclear protein; Cen
                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                       Caenorhabditis elegans.
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ZN_FING
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                                                                                                                                                                                                                                            F44E2.4.
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Local Similarity 40.9%;
nes 9; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SIMILARITY: Contains 1 GATA-type zinc finger.
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697 AA;
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78159 MW;
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                                                                                                                                                      Caenorhabditis
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HSSP; P01130; IAJJ;

WOXMPep; F4482.4; CE00182.

Pfam; PF00057; Idl recept_a.

SMART; SM00192; LDIa; 1.
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Bonfield J., Burton J., Connell M., Copsey T., Cooper J., Coulson A
Craxton M., Dear S., Du Z., Durbin R., Favello A., Fraser A.,
Fulton L., Gardner A., Green P., Hawkins T., Hillier L., Jier M.,
Johnston L., Jones M., Kershaw J., Kirsten J., Laisster N.,
Latreille P., Lightning J., Lloyd C., Mortimore B., O'Callaghan M.,
Parsons J., Percy C., Rifken L., Roopra A., Sounders D., Shownkeen
Sims M., Smaldon N., Smith A., Smith M., Sonnhammer E., Staden R.,
Sulston J., Thierry-Mieg J., Thomas K., Vaudin M., Vaughan K.,
Waterston R., Watson A., Weinstock L., Wilkinson-Sproat J.,
Materston R., Watson A., Weinstock L., Wilkinson-Sproat J.,
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                                                                                                                                                                                                                                                                                                           Hypothetical SEQUENCE 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      entities requires a license agreement (So or send an email to license@isb-sib.ch).
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                                                                       5 ILEGRIECVLSN 16
LLEGRISCVVSS
                                                                                                                                                                                                                                                                                                       al protein.
1609 AA;
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   888
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Pred. No.
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 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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2: pir2:*
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4: pir4:*
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T41352

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A18673

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42 35.9 67 2 S51401 42 35.9 231 2 T49830 42 35.9 255 2 A97877 42 35.9 260 2 C98084 42 35.9 283 2 S75226 42 35.9 322 2 D89785 42 35.9 365 2 JQ0198 42 35.9 418 2 A95168 42 35.9 418 2 B95197 42 35.9 418 2 B95197 42 35.9 418 2 B95146 42 35.9 418 2 H95208 42 35.9 418 2 T36944 42 35.9 482 2 T46944 42 35.9 514 2 T29652 42 35.9 562 2 E70399
562 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
T49830 A97877 C98084 C98084 C98085 D8975226 D897525 JQ0198 B95168 B95197 E95168 B95297 E95168 A95168 A95168 A95168 A95168 A95168 A95168 A95168 A95168 A95168 A95197 E95146 H95208

ALIGNMENTS

	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
A:Title: Cloning of the gene encoding the human erythropoletin receptor.	protein-tyrosine-p
	mypochiecter proce
A/NOTE: an insert compared to other Dubiness sense of the continues of the compared to the compared to sense of the compared to the compared t	propagate procean-c
A; NOTE: the authors translated the country that for restaue at as a	chrombosponani 3 -
A; Note: Bequence modified after extraction from NCBL backbone	therefore a
A; Kestades: I-I/; 36I-36/, LUEQUA , 39I-395; 504-506 < FEN-	nypothetical prote
A/MOLECULE Type: UNA	ISIL67, Cransposas
A; Accession: A55280	IS1167, transposas
A;Reference number: A55280; MUID:92147143; PMID:1664413	hypothetical prote
A; Title: Genomic organization of the human erythropoietin receptor gene.	degenerate transpo
	hypothetical prote
R; Penny, L.A.; Forget, B.G.	30S ribosomal prot
A; Residues: 1-508 < EHR >	protein PV100 [imp
A; Molecule type: mRNA	axonin 1 precursor
A;Accession: A53958	hypothetical prote
A53958; MUID:91372359;	probable transmemb
	ATP-dependent RNA
Exp. Hematol. 19, 973-977, 1991	F44E2.4 protein -
R;Ehrenman, K.; St. John, T.	probable transcrip
A; Note: sequence extracted from NCBI backbone (NCBIN:113293, NCBIP:113294)	cytochrome P450 Rv
	probable DNA-bindi
A;Cross-references: GB:S45332; NID:g255496; PIDN:AAB23271.1; PID:g255497	ribosomal protein
A; Residues: 1-508 < NOG>	ribosomal protein
A; Molecule type: DNA	thrombospondin 3 p
A;Accession: A49824	erythropoietin rec
A; Reference number: A49824; MUID: 92399733; PMID: 1668606	erythropoietin rec
A; Title: Cloning of the human erythropoietin receptor gene.	erythropoietin rec
Blood 78, 2548-2556, 1991	erythropoietin rec
R; Noguchi, C.T.; Bae, K.S.; Chin, K.; Wada, Y.; Schechter, A.N.; Hankine, W.D.	
A, Residues: 1-101, 'R', 103-188, 'RP', 191-243, 'E', 245-508 «WIN»	Description
A: Molecule type: mRNA: DNA	
A:Status: not compared with conceptual translation	
P-Persion Pagolago	
A/litte: The gene for the number effects problem receptor: analysis of the coulty sequence A.TBCFerence number AGOIGO. MITD 90304344: PMID:2163669;	cribution.
B1000 76, 24-30, 1990	esuit being printed,
R;Winkelmann, J.C.; Penny, L.A.; Deaven, L.L.; Forget, B.G.; Jenkins, R.B.	hance to have a
M60459; NID:g182244; PIDN:AAA52403.1; PID:g18	
A;Residues: 1-508 <jon></jon>	
A; Molecule type: mRNA	
A;Accession: A43799	
A; Reference number: A43799; MUID: 90304340; PMID: 2163696	
A.TH-le- Himman erythronoietin recentor: cloning expression and biologic characterizati	
RiJones, S.S.; D'Andrea, A.D.; Haines, L.L.; Wong, G.G.	
C;Accession: A43799; A60160; A49824; A53958; A55280; I52563	
C;Date: 12-Feb-1993 #sequence_revision 05-Apr-1995 #text_change 22-Jun-1999	
C; Species: Homo sapiens (man)	
erythropoietin receptor precursor - human	
ZUHUR	

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A;Cross-references: GDB:125242; OMIM:133171

A;Map position: 19p13.3-19p13.2

A;Introns: 39/1; 84/2; 143/1; 195/3; 247/1; 276/2; 305/3

A;Introns: 39/1; 84/2; 143/1; 195/3; 247/1; 276/2; 305/3

C;Superfamily: erythropoietin receptor; cytokine receptor homology
C;Keywords: alternative splicing; cytokine receptor; glycoprotein; transmembrane protein
F;1-24/Domain: signal sequence #status predicted <SIG>
F;25-508/Product: erythropoietin receptor #status predicted <MAT>
F;52-250/Domain: cytokine receptor bomology <CRS>
F;52-239/Domain: cytokine receptor bomology <CRS>
F;23-237/Region: WSXWS motif
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C;Keywords: cytob
F;52-238/Domain:
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A; Residues: 1-265 < KUR>
A; Residues: C.; Chretien, S.; Lemarchandel, V.; Mayeux, P.; Romeo, P.
J. Biol. Chem. 266, 6952-6956, 1991
J. Title: Spleen focus-forming virus long terminal repeat insertional activation of
A; Reference number: 149653; MUID:91201346; PMID:1849897
A; Accession: 149653
erythropoietin receptor precursor, membrane-bound form - C;Species: Mus musculus (house mouse)
C;Date: 28-Sep-1990 #sequence_revision 05-Apr-1995 #text_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Title: Characterization of murine erythropoietin receptor genes. A;Reference number: S13249; MUID:91080149; PMID:2175360 A;Accession: S14081
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C;Accession: S14081; I49653
R;Kuramochi, S.; Ikawa, Y.; Todokoro,
J. Mol. Biol. 216, 567-575, 1990
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A; Residues: 1-24 < RES>
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C;Genetics:
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A;Residues: 1-96 <RES>
                                                                                       A32385
                                                                                                                 RESULT 3
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|Species: Mus musculus (house mouse)
|Jate: 18-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Superfamily: erythropoietin receptor; cytokine receptor homology; Keywords: cytokine receptor; transmembrane protein
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                          cytokine receptor homology <CRS>
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#sequence_revision 05-Apr-1995 #text_change 22-Jun-1999
                                                                                                                                                                                                                                                                                                                                         93.2%;
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                                                                                                                                                                                                                                                                                                               1; Mismatches
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Pred. No. 2.3e-09;
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C;Superfamily: erythropoletin receptor; cytokine receptor homology C;Keywords: cytokine receptor; glycoprotein; transmembrane protein F;1-24/Domain: signal sequence #status predicted <SIG> F;25-507/Product: erythropoletin receptor #status predicted <WAT> F;25-249/Domain: extracellular #status predicted <EXT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Accession: A46713
R;Masuda, S.; Nagao, M.; Takahata, K.; Konishi, Y.; Gallyas Jr J. Biol. Chem. 268, 11208-11216, 1993
A;Title: Functional erythropoietin receptor of the cells with A;Reference number: A46713; MUID:93266574; PMID:7684373
A;Accession: A46713
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c;Superfamily: erythropoietin receptor; cytokine receptor homology
C;Keywords: alternative splicing; cytokine receptor; glycoprotein;
F;1-24/Domain: signal sequence #status predicted <SIG>F;1-24/Domain: signal sequence #status predicted <SIG>F;25-507/Product: erythropoietin receptor #status predicted <MAT>F;25-249/Domain: extracellular #status predicted <EXT>F;25-238/Domain: cytokine receptor homology CCRS>F;25-238/Domain: cytokine receptor #status predicted complexity cytokine receptor #status predicted 
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A;Residues: 1-507 <KUR>
A;Cross-references: EMBL:X53081; NID:g50861; PIDN:CAA37248.1; PID:g50862
A;Experimental source: murine erythroleukemia K-1 cells
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A;Cross-references: GB:J04843; NID:g193090; PIDN:AAA37571.1; PID:g309219
A;Cross-references: murine erythroleukemia (MEL) cells, subclone 745
A;Experimental source: murine erythroleukemia (MEL) cells, subclone 745
R;Kuramochi, S.; Itawa, Y.; Todokoro, K.
J. Mol. Biol. 216, 567-575, 1990
A;Title: Characterization of murine erythropoietin receptor genes.
A;Reference number: S13249; MUID:91080149; PMID:2175360
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A;Title: Expression cloning of the murine erythropoietin receptor. A;Reference number: A32385; MUID:89195238; PMID:2539263
A;Accession: A32385
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Mol. Cell. Biol. 11, 5527-5533, 1991
A;Title: Unregulated expression of the erythropoietin receptor gene caused by insertion of A;Reference number: A41686; MUID:92017832; PMID:1656233
                                                                                                                                                                                                                                                                                                            A;Cross-references: GB:D13566; NID:g286209; PIDN:BAA02761.1; PID:g286210
A;Experimental source: PC12 and erythroid cells
A;Note: sequence extracted from NCBI backbone (NCBIN:132811, NCBIP:132813)
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A; Residues: 1-507 <HIN>
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R;Hino, M.; Tojo, A.; Misawa, Y.; Morii, H.; Takaku, F.; Shibuya, M.
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A;Molecule type: mRNA
A;Residues: 1-507 <MAS>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;Species: Rattus norvegicus (Norway rat)
;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 16-Jun-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21;
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cytokine
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receptor homology
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Pred. No. 4.6e-09;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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e F5-5
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RESULT 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Species: Homo sapiens (man)
C;Date: 03-Nov-1995 #sequence revision 03-Nov-1995 #text_change 21-Jan-2000
C;Accession: A57121; T08830
R;Adolph, K.W.; Long, G.L.; Winfield, S.; Ginns, E.I.; Bornstein, P.
Genomics 27, 329-336, 1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A,Description: mediates cell-to-matrix and cell-to-cell interactions C;Superfamily: thrombospondin 3; EGF homology F;374-412/Domain: EGF homology <EGF>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Map position: 1q21-1q23
A;Introns: 27/1; 96/1; 181/3; 216/1; 225/1; 256/1; 270/1; 319/3; 366/3; 392/3; 443/3; 48
C;Function:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Title: Structure and organization of the human thrombospondin 3 gene (THBS3). A;Reference number: A57121; MUID:96044440; PMID:7558000 A;Accession: A57121
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                                                                                                    Yeast 8, 949-959, 1992

A;Title: Structure and expression of the ABF1-regulated ribosomal protein S33

A;Reference number: S30005; MUID:93127729; PMID:1481571

A;Accession: S30006
                                                                                                                                                                                                                                                               ribosomal protein S28.e - yeast (Kluyveromyces marxianus) N;Alternate names: ribosomal protein YS33 C;Species: Kluyveromyces marxianus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Cross-references: EMBL:AF023268; NID:g2564910; PIDN:AAC51818.1; C;Genetics:
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A;Molecule type: DNA
A;Residues: 1-736 <WIN>
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A; Residues: 1-956 < ADO>
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                                                                                                                                                                                          C;Accession: S30006
R;Hoekstra, R.; Ferreira, P.M.; Bootsman, T.C.; Mager, W.H.; Planta, R.J.
                                                                                                                                                                                                                      C;Date: 07-May-1993 #sequence_revision 07-May-1993 #text_change 07-May-1999 C;Accession: S30006
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A;Cross-references: GDB:409953; OMIM:188062
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                                        A;Cross-references: GB:S52656
                                                              A; Molecule type: DNA
A; Residues: 1-67 < HOE>
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Pred. No. 41;
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Pred. No. 4.6e-09;
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A;Gene: S33
C;Superfamily: rat ribosomal protein S28
C;Keywords: protein biosynthesis; riboso
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A;Accession: S30005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N,Alternate names: ribosomal protein YS33
C;Species: Kluyveromyces marxianus var. lactis, Candida sphaerica
C;Date: 07-May-1993 #sequence_revision 07-May-1993 #text_change 07-May-1999
C;Accession: S30005
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                                                                                                                                                                                                                                                                                                                                   A;Title: Genome sequence of a human tumorigenic poxvirus: Prediction of specific host A;Reference number: Z20876; MUID:96325459; PMID:8670425
                                                                                                                                                                                                                                                                                                                                                                            R; Senkevich, T.G.; Bugert,
Science 273, 813-816, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         probable DNA-binding protein 46L - Molluscum contagiosum virus N,Alternate names: MCO46L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Cross-references:
C; Genetics:
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A; Residues: 1-67 < HOE>
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                                                                                                                                                         C; Keywords:
                                                                                                                                                                                                                                                          A; Molecule type: DNA
A; Residues: 1-288 <SEN>
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                                                                                                                                                                                              A; Note: MC046L
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                                                                                                                                                                                                                                                                                                                    A;Accession: T30648
                                                                                                                                                                                                                                                                                                                                                                                                                 Species: Molluscum contagiosum virus 1
,Date: 05-Nov-1999 #sequence_revision 05-Nov-1999 #text_change 27-Oct-2003
,Accession: T30648
                                                                                                                                                                                                                   ;Genetics
                                                                                                                                                                                                                                                                                             Status: preliminary; translated
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Best Local
                                                                                                                                                                         Superfamily: DNA-binding phosphoprotein, vaccinia I3L type
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les 9; Conserv
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les 8; Conserv
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133
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                                                                                                                                                       DNA binding
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ECIYTNLRGGTR 144
                                   ECVLSNLRGRTR 22
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                                                                              Conservative
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42.9%; Pred. No. 3.
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Pred. No.
                                                                            Score 46; DB 2;
Pred. No. 17;
2; Mismatches
                                                                                                                                                                                                                                                                                             from GB/EMBL/DDBJ
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C;Species: Mycobacterium tuberculosis
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 21-Jul-2000
C;Accession: H70729
R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Squares, Connor, R.; Davies, R.; Davies, R.; Davies, R.; Davies, R.; Connor, R.; Davies, R.; Davies, R.; Feltwell, T.; Gentles, S.; Squares, R.; Connor, R.; Davies, R.; Davies, R.; Feltwell, T.; Gentles, S.; Squares, R.; Squar
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A;Molecule type: DNA
A;Residues: 1-710 <BAR>
A;Cross-references: EMBL:AL117389; PIDN:CAB55767.1; GSPDB:GN00068;
A;Experimental source: strain 972h-; cosmid c4F11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     probable transcription regulator - fission yeast (Schizosaccharomyces pombe) C;Species: Schizosaccharomyces pombe C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 21-Jan-2000 C;Accession: T41352; T38826 C;Accession: T41352; T38826 R;Barrell, B.G.; Rajandream, M.A.; Quail, M.; Harris, D.; Lyne, M. submitted to the EMBL Data Library, June 1999
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A;Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G. A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complete A;Reference number: A70500; MUID:98295987; PMID:9634230
A;Accession: H70729
A;Status: preliminary; nucleic acid sequence not shown; translation not shown A;Molecule type: DNA
A;Residues: 1-489 <COL>
                                                                                                                                                                                                    F44E2.4 protein - Caenorhabditis elegans
C;Species: Caenorhabditis elegans
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 07-Aug-1998
C;Accession: S44821
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                                A; Reference number: S44816
A; Accession: S44821
                                                                                                     submitted to the EMBL Data Library, September 1993 A_iDescription: Sequence of the C. elegans cosmid F
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A;Accession: T41352
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A;Status:
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A;Map position: 3
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Pred. No. 29;
4; Mismatches
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C;Genetics:
A;Gene: PA1S
C;Superfamil
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A;Residues: 1-1609 <AND>
A;Residues: 1-1609 <AND>
A;Cross-references: EMBL:L23646; NID:g388595; PID:g388601
A;Cross-references: EMBL:L23646; NID:g388595; PID:g388601
C;Genetics:
C;Genetics:
A;Introns: 107/2; 173/2; 228/3; 594/3; 1165/2; 1216/3; 1231/3; 1258/2; 1300/2; 1321/3; 1:
C;Superfamily: LDL receptor ligand-binding repeat homology
F;11-45/Domain: LDL receptor ligand-binding repeat homology <LDL>
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D83406
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S42639
                                                                                                                                                                                                                                        A;Title: Complete genome sequence of Pseudomonas aeruginosa
A;Reference number: A82950; MUID:20437337; PMID:10984043
A;Accession: D83406
                                                                                                                                                                                                                                                                                                     R;Stover, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, adman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, .; Lory, S.; Olson, M.V.
Nature 406, 959-964, 2000
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                                                                                                                                          A;Cross-references: GB:AE00461
A;Experimental source: strain
                                                                                                                                                                                A; Molecule type: DNA
A; Residues: 1-316 < STO>
                                                                                                                                                                                                                                                                                                                                                                                   C;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 28-Jul-2003 C;Accession: D83406
                                                                                                                                                                                                                                                                                                                                                                                                                         probable transmembrane sensor PA1911 [imported] - C;Species: Pseudomonas aeruginosa
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                                                                               Gene: PA1911
Superfamily:
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    Conservative
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                                                                                                                                                             GB:AE004617; GB:AE004091; NID:g9947901; PIDN:AAG05299.1; GSPDB:GN001:
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                  38.5%;
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Pred. No. 44;
4; Mismatches
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Pred. No.
                  Score 45;
Pred. No.
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    Mismatches
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le+02;
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                                         Length 316;
                                                                                 component
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A.; Larbig,
    Gaps
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K.; Lim,
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A;Molecule type: protein
A;Residues: 29-49;51-80;84-95;100-117;120-128;130-141;143-176;243-254;256-296;303-336;33
A;Residues: 29-49;51-80;84-95;100-117;120-128;130-141;143-176;243-254;256-296;303-336;33
R;Gigger R.J.; Vogt, L.; Zuellig, R.A.; Rader, C.; Henehan-Beatty, A.; Wolfer, D.P.; Sor Bur. J. Biochem. 227, 617-628, 1995
Bur. J. Biochem. 227, 617-628, 1995
A;Title: The gene of chicken axonin-1. Complete structure and analysis of the promoter.
A;Reference number: S69332; MUID:95172044; PMID:7867620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N;Alternate names: neural cell adhesion molecule AxCAM C;Species: Gallus gallus (chicken)
C;Species: Gallus gallus (chicken)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 21-Jan-2000
C;Accession: S22383; S34107; S69332; S22128
R;Zuellig, R.A.; Rader, C.; Schroeder, A.; Kalousek, M.B.; von Bohlen und Halbach, F.; Bur. J. Biochem. 204, 453-463, 1992
A;Title: The axonally secreted cell adhesion molecule, axonin-1. Primary structure, imm A;Reference number: S22383; MUID:92174898; PMID:1311675
A;Accession: S22383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hypothetical protein Rv2733c - Mycobacterium tuberculosis (strain H37RV)

(;Species: Mycobacterium tuberculosis

C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 21-Jul-2000

C;Accession: D70506

R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Gordon, S.; Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Holroyd, S. Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S. Nature 393, S37-S44, 1998

Nature 393, S37-S44, 1998

A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complete genome A;Reference number: A70500; MUID:98295987; PMID:9634230
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, June 1994 C;Superfamily: contactin; fibronectin type III repeat homology; immunoglobulin homology c;Keywords: cell adhesion F;1-23/Domain: signal sequence #status predicted <SIG>F;24-1036/Product: axonin 1 #status predicted <MAT>F;336-392/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                             A;Status: preliminary; nucleic acid sequence not shown; translation not shown A;Molecule type: DNA A;Residues: 1002-1036 <GIG>
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A; Residues: 1-1036 < ZUE1>
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C;Superfamily: conserved hypothetical protein b0835
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A; Residues: 1-512 < COL>
                                                                                                                                                                                                                                         A; Cross-references: EMBL: X79607
                                                                                                                                                                                                                                                                                                                                                                                                 A; Accession: S69332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Accession: S34107
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cross-references: EMBL:X63101; NID:g62852; PIDN:CAA44815.1; PID:g62853
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Local Similarity 42.3%;
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Pred. No. 44;
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Query Match

38.5%;

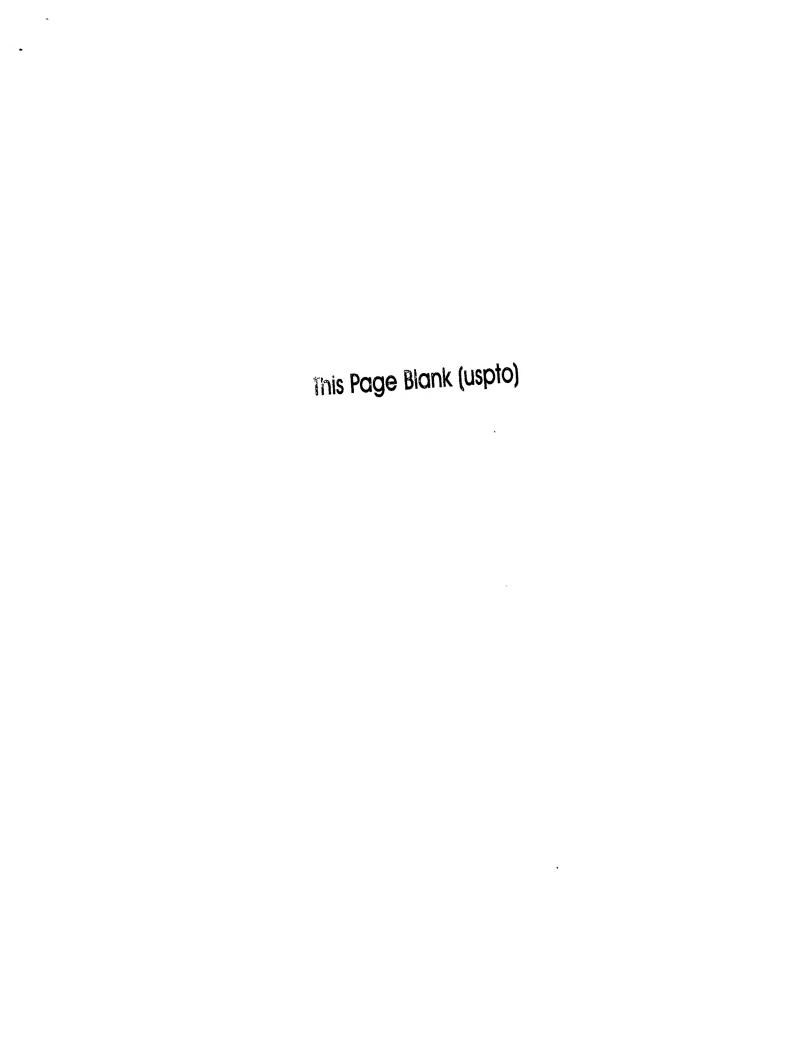
Score 45;

DB 2;

Length 1036;

0;

Search completed: May 6, 2004, 12:47:27 Job time: 22 secs



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Title:
Perfect score:
Sequence:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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A Geneseq_29Jan04:*
1: geneseqp1980s:*
2: geneseqp1990s:*
3: geneseqp2900s:*
4: geneseqp2001s:*
5: geneseqp2003as:*
6: geneseqp2003bs:*
7: geneseqp2003bs:*
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120.344 Million cell updates/sec
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Gapop 10.0 , Gapext 0.5
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117
                                                                                                                                                                                                                                                                                                                                                                               1586107 seqs, 282547505 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              QRVEILEGRTECVLSNLRGRTRY 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	. 10	9	8	7	Ð	տ	4	ω	N	ב	Result No.
52	102	109	109	109	109	111	117	117	117	117	117	117	117	117	117	117	117	117	117	117	117	117	117	117	Score
44.4	87.2	93.2	93.2	93.2	93.2	94.9	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	Query Match I
1026	507	507	507	507	265	508	508	508	508	508	508	503	488	458	458	458	438	438	225	211	30	23	23	23	Length I
σı	ນ	N	N	N	N	7	ຫ	N	N	N	N	ω	N	7	7	7	ω	ω	w	w	N	σ	տ	N	BB
AAU80379	AAR50327	AAR69502	AAR47517	AAR06511	AAR50326	ADE28677	ABB09173	AAR69503	AAR70032	AAR47518	AAR06512	AAB13012	AAW08349	ADE28675	ADE28673	ADE28671	AAY44623	AAY44622	AAB21685	AAB21686	AAR98938	ABP72837	AAU78011	AAY39411	ID
Aau80379 Human BIG	Mouse	Aar69502 Mouse ery	MEL	EPO	٠.	Human	Human	Human	Human	Aar47518 Human EPO	EPO r	Aab13012 Q-tagged		Ade28675 Human NOV	Ade28673 Human NOV	Ade28671 Human NOV	R154C	Aay44622 Truncated	Aab21685 Human mat	Aab21686 Human mat	Aar98938 Synthetic	Abp72837 Erythropo		39411 Human e	Description

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41.9	41.9	41.9		41.9		41.9	41.9	41.9	41.9	41.9	41.9	41.9	41.9	41.9	41.9	41.9		41.9	
470	470	458	458	456	456	451	451	413	413	373	373	341	321	286	221	199	49	49	1
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Abu08371	Aab66083	Abg76512	Aay02135	Abo32693	Aab66033	Abo32717	Aab66063	Aae13244	Aab84220	Abo32694	Aab66034	Aab97378	Aay02136	Abp43791	Abp65639	Abg13674	Aae13249	Aab84225	
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ALIGNMENTS

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This sequence represents a peptide derived from the extracellular activation sequence of the human crythropoletin (BPO) receptor. Activation sequences are involved in modulation of receptor responses and are separate from the ligand binding site. Activation sequences are important in two distinct ways: in the modulation of receptor internalisation; and/or in the modulation of activation of the receptor use of peptides corresponding to the activation sequence of a receptor can retard or inhibit receptor internalisation, thereby increasing or stabilising the steady-state number of active receptors on the cell surface. This has the effect of increasing signalling per unit of ligand. In addition, the receptor is classed as type 2 cell surface receptor, meaning that such peptides can actually replace the requirement for the ligand, causing ligand-independent activation. This activation is probably brought about by a dimerisation mechanism in which one peptide	ptors	WO9942127-A2. 26-AUG-1999. 23-FEB-1999; 99WO-US003910. 24-FEB-1998; 98US-00028938. (RECE-) RECEPTRON INC.	30-NOV-1999 (first entry) Human erythropoietin receptor-derived activation peptide. Erythropoietin; EPO; receptor; activation; internalisation. Synthetic. Homo sapiens.	RESULT 1 AAY39411 standard; peptide; 23 AA. ID AAY39411 standard; peptide; 23 AA. XX AAY39411;

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This invention corresponds to a novel composition comprising any of 34 CC peptides consisting of activation sequences from the extracellular CC domains of cell-surface receptors. These peptides were identified by CC homology searching with a peptide sequence from the MHC class I alpha I CC receptor. Activation sequences are involved in the internalisation of CC receptors and so these peptides may be used to modulate the CC internalisation and or activation of these receptors. The peptides of the CC invention may also be used to increase effect of ligand signalling in CC type 2 diabetes and can replace the ligand normally required for CC activation. This is useful when the ligand is a hormone and is difficult to obtain. The peptides may also interast synergistically with the ligand or function as antagonists of receptor signalling. The peptides can also be used in drug screening to identify compounds that modulate receptor conternalisation or function as ligand replacements. Use of these peptides can increase the effect of therapeutic hormones (optionally codaminate receptors at least 50% and are specific for the receptors from CC which they derive. The present sequence represents the human content of the components of the receptor activation sequence oligopeptide of the invention contents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                          Composition comprising peptides from receptor extracellular domains, useful for e.g. modulating receptor internalization and activation such as increasing insulin activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                subsequent activation that occurs when two receptors bind one ligand. This peptide could be used to treat disorders involving an inadequate inappropriate response from its corresponding receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Olsson L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Erythropoietin receptor; antidiabetic; activation sequence; ligand; receptor internalisation; type 2 diabetes; drug screening; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-MAR-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAU78011 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RECE-) RECEPTRON INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2002-224899/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            erythropoeitin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Naranda T;
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                                                                                                                                                                                                                                                                                                                                                                                       11; 48pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                96US-00612999.
96US-00701382.
97US-00788820.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          receptor activation domain oligopeptide
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Pred. No. 3.6e-11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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Best Local :
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                                              The present sequence is that of an erythropoietin (EPO) mimetic peptide which has no homology to natural EPO but has similar biological properties in that it activates the EPO receptor acting as an agonist. The EPO mimetic peptide can be included in novel fusion proteins of the invention also including a transferrin (Tf. see ABP72820) molety engineered to extend the serum half-life or bioavailability. The EPO mimetic peptide can be fused to the N- or C-terminus of Tf. or inserted into, or used to replace part of Tf. such that the Tf acquires EPO activity. Modified Tf fusion proteins of the invention can be used in the diagnosis, prognesis, prevention and/or treatment of diseases and/or disorders of the endocrine, nervous, immune, respiratory, cardiovascular, reproductive and digestive systems, diseases and/or disorders relating to the blood or to cell proliferation, inflammatory conditions and infectious diseases, or to deliver a therapeutic agent to a cell or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                    New fusion protein, useful in the diagnosis and treatment of diseases or disorders relating to the respiratory, cardiovascular and digestive systems, comprises a transferrin protein fused to a therapeutic protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fungicide; hepatotropic; antirheumatic; antiarthritic; antigout; tranquillizer; vulnerary; antidiabetic; nephrotropic; antipyretic; gastrointestinal; gene therapy; transgenic animal; erythropoietin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Transferrin; neuroprotective; cerebroprotective; vasotropic; antiparkinsonian; nootropic; anti-HIV; antiasthmatic; antiallergic; cytostatic; immunosuppressive; antiatherosclerotic; cardiant;
Sequence
                                                                                                                                                                                                                                                                                                                                   Example 9; Page 255; 298pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-332916/31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-AUG-2001; 2001US-0315745P 30-NOV-2001; 2001US-0334059P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-AUG-2002; 2002WO-US027637
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gynaecological; immunostimulant; antianaemic; haemostatic;
antiinflammatory; dermatological; antibacterial; virucide; antiparasitic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Erythropoietin mimetic peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-AUG-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mimetic; agonist.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (BIOR-) BIOREXIS PHARM CORP
                                    the blood-brain barrier
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 ORVEILEGRIECVLSNLRGRIRY 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard; peptide;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 117; DB 5; 100.0%; Pred. No. 3.6e-11;
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RESULT 4
AAR98938
ID AAR9
RESULT 5
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ID AAB2
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                                                                                                                                                                                  Matches
                                                                                                                                                                                             Query Match
Best Local
                                                                                                                                                                                                                                                           contg. the antibody may be used in the diagnosis and treatment of patients having disorders associated with low red blood cell levels, e. anaemia. The antibodies are also useful in methods and kits for detecti EPORs in biological in biological samples
                                                                                                                                                                                                                                                                                                                                         AAR89960-R89965 and AAR98936-R98939 are overlapping, synthetic human erythropoietin receptor (shuEPOR) peptides which span residues 1 to 244 of the human EPOR. The peptides are used to map the EPOR binding epitope of an EPOR monoclonal antibody which binds to EPORs and stimulates
                                                                                                                                                                                                                                                                                                                                                                                                                                    Monoclonal antibodies stimulating an erythropoietin receptor - useful in diagnosis and treatment of patients having disorders associated with low red blood cell levels, e.g. anaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR98938;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR98938 standard;
                                                                                                                                                                                                                                     Sequence 30 AA;
                                                                                                                                                                                                                                                                                                                erythropoiesis by stimulating the proliferation and/or differentiation of erythroid progenitor cells to erythrocytes. Pharmaceutical compsns.
                                                                                                                                                                                                                                                                                                                                                                                                            Example 6; Page 34; 61pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-117004/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      erythropoiesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic human erythropoietin receptor peptide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-SEP-1996
               21-DEC-2000
                                       AAB21686;
                                                                AAB21686 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-JUL-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Monoclonal antibody; erythropoietin receptor; diagnosis; anaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (AMGE-) AMGEN INC.
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                                                                                                                                                                                              Similarity
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                                                                                                                               QRVEILEGRIECVLSNLRGRIRY 23
                                                                                                                                            QRVEILEGRIECVLSNLRGRIRY 23
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                                                                                                                                                                                  Conservative
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               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    erythrocyte production; epitope mapping.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; --
                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                            English.
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                                                                211 AA
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                                                                                                                                                                                             Score 117; DB 2;
Pred. No. 4.9e-11;
                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              . 3.6e-11;
ches 0;
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                                                                                                                                                                                                         Length 30;
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                                                                                                                                                                                                                                                                                           RESULT 6
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Screening for ligand analogs and agents which modulate ligand-receptor binding, comprises adding a test ligand to a non-naturally occurring constraint ecceptor analog.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human mature erythropoietin receptor BPOR extracellular domain
                                     11-FEB-1999;
29-APR-1999;
                                                                                                                      WO200047612-A2
                                                                                                                                             Homo sapiens.
                                                                                                                                                                                           Human mature erythropoietin receptor EPOR extracellular domain
                                                                                                                                                                                                                     21-DEC-2000
                                                                                                                                                                                                                                             AAB21685;
                                                                                                                                                                                                                                                                   AAB21685 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 211 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        analog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Fig 8; 82pp; English.
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29-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ligand; cell surface receptor; erythropoietin; EPOR; human;
                                                                        11-FEB-2000; 2000WO-US003665
                                                                                                                                                                    Ligand; cell surface receptor; erythropoietin; EPOR; human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (XENC-) XENCOR INC.
                                                                                               17-AUG-2000.
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                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                 QRVEILEGRTECVLSNLRGRTRY 23
                                                                                                                                                                                                                                                                                                                                 QRVEILEGRTECVLSNLRGRTRY 183
                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                     (first entry)
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99US-0131674P.
                                     99US-0120009P
99US-0131674P
                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 117; DB 3; 100.0%; Pred. No. 4.7e-10;
                                                                                                                                                                                                                                                                     225
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(XENC-) XENCOR INC

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RESULT 7
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Best Local :
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        The present sequence is truncated human EpoR (erythropoietin receptor). Transfection of constitutively active EpoR (t449) by electroporation into a cytokine-dependent cell line supports cell population expansion in the absence of exogenous cytokines. Mutant human EpoR is used in treatment of disorders related to inadequate EpoR signalling. The transfected cells
                                                                                                                                                                                                                                                                                                                                                                      Truncated human EpoR; erythropoietin receptor; hypersensitive EpoR(t439); mutant human EpoR; EpoR signalling; cancer; infectious disease; HIV; sickle cell anaemia; cytostatic; antimicrobial; antiviral;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to a method for screening for a ligand analog, comprising adding a candidate ligand to a non-naturally occurring cell surface receptor analog e.g. erythropoietin receptor (EPOR), and determining the binding of the ligand to the analog. The present sequence is a mature human erythropoietin receptor (EPOR) extracellular domain.
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binding, c
                                                                                 Example
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                                                                                                                                                                                                                                        25-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                       Truncated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAY44622;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY44622 standard; protein; 438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example
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                                                                                                                                                                                                                                                                                                                                                              immunostimulant;
                                                                                                                                                                                                      (HEMO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence may
                                                                                                                                                       2000-136979/12.
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                                                                                                                    free defined medium useful
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             μ
                                                                                                                                                                                                      HEMOSOL INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               receptor analog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dahiyat
                                                                                 6
                                                                                                                                            AAZ49634.
                                                                                                                                                                               Matthews KE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     QRVEILEGRIECVLSNLRGRIRY 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            QRVEILEGRTECVLSNLRGRTRY 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  225 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         analog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       y for ligand analogs and comprises adding a test
                                                                                                        production
                                                                                                                                                                                                                                                                                                                                                                                                                       human
                                                                                Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fig 8; 82pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                            98CA-02241576.
99CA-02260332.
                                                                                                                                                                                                                                                                99WO-CA000606.
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                                                                                                                                                                                                                                                                                                                                                                                                                       EpoR (t439).
                                                                                                                                                                                                                                                                                                                                                               anti-anaemic
                                                                               61pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     be used in the present invention as a cell surface
                                                                                                         ę,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
therapy to
                                                                                                        hemoglobin
                                                                                                                                                                               Mueller
                                                                                English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 117; DB 3
Pred. No. 5e-10;
                                                                                                                                                                               SG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                    for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          agents which modulate ligand-receptor ligand to a non-naturally occurring cell
                                                                                                                     the
                                                                                                                     efficient
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                                                                                                                     culture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            225;
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                                                                                                                     stem
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                                                                                                                     cells
                       of.
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QRVEILEGRTECVLSNLRGRTRY 23

Query Match Best Local S Matches 23

Similarity 23; Conser

100.0%; ilarity 100.0%; Conservative 0;

0

Mismatches

0

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0

Length 438;

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RESULT 8
AAY44623
ID AAY4
AC A
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Best Local S
Matches 23
                                                                                                                                                                                  The present sequence is R154C truncated human EpoR (erythropoietin receptor). Transfection of constitutively active EpoR (e439; R154C) electroporation into a cytokine-dependent cell line supports cell population expansion in the absence of exogenous cytokines. Mutant EpoR is used in treatment of disorders related to inadequate EpoR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HIV),
                                                                                                               EpoR is used in treatment of disorders related to inadequate EpoR signalling. The transfected cells may also used in gene therapy to cancer, infectious diseases (e.g. HIV), sickle cell anaemia, and conditions related to abnormal expression of erythropoietin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Truncated human EpoR; erythropoietin receptor; hypersensitive EpoR(t439); mutant human EpoR; EpoR signalling; cancer; infectious disease; HIV; sickle cell anaemia; cytostatic; antimicrobial; antiviral; immunostimulant; anti-anaemic.
                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                   used
                                                                                                                                                                                                                                                                                                                                                                                                            Serum
                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bell D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-JUN-1998;
25-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9967360-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R154C truncated human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-APR-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 438
                                                                                                                                                                                                                                                                                                                                    Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (-OMBH)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            v), sickle cell anaemia,
erythropoietin
                                                                                                                                                                                                                                                                                                                                                                                   for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000-136979/12
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                                                                                                                                                                                                                                                                                                                                                                                 free defined medium useful for production of hemoglobin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 QRVEILEGRTECVLSNLRGRTRY 23
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                                                                       438
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matthews KE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                  Fig 10; 61pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A,
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99CA-02260332.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "Wild type Arg substituted by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EpoR (t439).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mueller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and conditions related to abnormal expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
Score 117; DB 3;
Pred. No. 1.1e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 117; DB 3;
Pred. No. 1.1e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     438;
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QRVEILEGRIECVLSNLRGRIRY 216

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RESULT 9
ADE28671
ADE
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17-MAY-2002;
28-MAY-2002;
28-MAY-2002;
28-MAY-2002;
29-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-DEC-2001;
07-DEC-2001;
12-DEC-2001;
17-DEC-2001;
17-DEC-2001;
20-DEC-2001;
27-DEC-2001;
31-DEC-2001;
17-APR-2002;
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                                                                                                                                                                                                                                        Alsobrook JP, Alvarez E, Anderson DW, Baron M, Boldog FL; Burgess CE, Casman SJ, Chapoval A, Dhanabal M, Edinger SR, Ellerman K, Ettenberg S, Gangolli EA, Gerlach VL, Gorman I Grosse WM, Guo X, Hackett C, Ji W, Kekuda R, Khramtsov NN Lepley DM, Li L, Macdougall JR, Malyankar UM, Mazur A, Mezes PS, Miller CE, Miller I, Mishra VS, Padigaru M, Pat Pena CEA, Peyman JA, Rastelli I, Rieger DK, Shenoy SG, St Smithson G, Starling G, Spytek KA, Stone DJ, Tchernev VT, Vernet CAM, Zerhusen BD, Zhong M;
                                                    New isolated NOVX polypeptides and polynucleotides, useful for preventing, diagnosing or treating NOVX-associated disorders, e.g. osteoarthritis, obesity, atherosclerosis, cancer, Parkinson's disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human
                                                 preventing, diagnosing or treating NOVX-as osteoarthritis, obesity, atherosclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-NOV-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CURA-)
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                             or intections
                                                                                                                                                                   ADE28670
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2001US-03413495P.
2001US-0341477P.
2001US-0341540P.
2001US-0344297P.
2001US-0344297P.
2001US-0344293P.
2002US-0380981P.
2002US-0380981P.
2002US-0381495P.
2002US-0381495P.
2002US-0381495P.
2002US-0381495P.
2002US-0383744P.
2002US-0383744P.
2002US-0383744P.
2002US-0383744P.
2002US-0383744P.
2002US-0383743P.
2002US-0383743P.
2002US-0383747971.
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Edinger SR,
                                                                                                                                                                                                                                                                                                   Patturajan
, Shimkets I
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Z
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RESULT 10
ADE28673
ID ADE286
XX ADE28
XX ADE28
XX ADE28
XX NOVX;
XW ANTL:
XW NOVX;
XW ANTL:
XW NOUTC
XW ANTL:
XW OSTEC
XX ISAN
XX
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Best Local S
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05-NOV-2001;

05-DEC-2001;

07-DEC-2001;

12-DEC-2001;

17-DEC-2001;

17-DEC-2001;

20-DEC-2001;

21-DEC-2001;

31-DEC-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NOVX; antidiabetic; anorectic; cardiant; hypotensive; antiarteriosclerotic; virucide; antibacterial; fungicide; protozoacide; nootropic; neuroprotective; antiparkinsonian; anticonvulsant; osteopathic; antiarthritic; antiinflammatory; dermatological; antiasthmatic; antilipaemic; metabolic; diabetes; obesity; infectious; anorexia; cancer; cardiovascular; hypertension; atherosclerosis; neurodegenerative; Alzheimer's disease; Parkinson's; epilepsy; immune; osteoarthritis; haemopoietic; inflammatory skin; asthma; dyslipidaemia; neurogenesis; cell differentiation; proliferation; haemopoiesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 458 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human NOV15b protein - SEQ ID 50.
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                                                                                                                                                                                                                                                                                                              05-NOV-2002;
                                                                                                                                                                                                                                                                                                                                                                          15-MAY-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     wound healing; angiogenesis; gene therapy; chromosome
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2001US-0338626P.

2001US-0336600P.

2001US-0341346P.

2001US-0341477P.

2001US-0341477P.

2001US-0341540P.

2001US-0342592P.

2001US-0344297P.

2001US-0344903P.
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Pred. No. 1.1e-09;
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15-MAY-2002;
17-MAY-2002;
28-MAY-2002;
28-MAY-2002;
29-MAY-2002;
29-MAY-2002;
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Lepley DM,
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Burgess CE, C
Ellerman K, E
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Smithson G,
                                                                                                                                                                                                                                                                                                                                  disease and epilepsy, immune disorders e.g. osteoarthritis, haemopoietic disorders, inflammatory skin disorders, asthma and dyslipidaemia. Furthermore, the nucleic acids and polypeptides may also be used to identify molecules that modulate or inhibit neurogenesis, cell differentiation and proliferation, haemopoiesis, wound healing and angiogenesis, as well as in gene therapy. Finally, the nucleic acids may be used as hybridisation probes, in chromosome mapping, tissue typing, preventive medicine and pharmacogenomics. The current sequence is that the human NOV protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       anticonvulsant, osteopathic, antiarthritic, antiinflammatory, dermatological, antiasthmatic and antilpaemic activities. The polypeptides, nucleic acid molecules and antibodies may be useful freating or diagnosing diseases including metabolic disorders such diabetes and obesity, infectious diseases, anorexia, cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a novel isolated NOVX polypeptide. The polypeptide of the invention demonstrates, antidiabetic, anorectic, cardiant, hypotensive, antiarteriosolerotic, virucide, antibacterial, fungicide, protozoacide, nootropic, neuroprotective, antiparkinsonian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New isolated NOVX polypeptides and polynucleotides, useful for preventing, diagnosing or treating NOVX-associated disorders, e.g. psteparthritis, obesity, atherosclerosis, cancer, Parkinson's disease,
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                   29-JAN-2004
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DB; ADE28672.
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, Casman SJ, Chapoval A, Dhanabal M, Edinger SR,
Cattenberg S, Gangolli EA, Gerlach VL, Gorman L;
Guo X, Hackett C, Ji W, Kekuda R, Khramtsov NV;
Li L, Macdougall JR, Malyankar UM, Mazur A, Mcqn
Miller CB, Millet I, Mishra VS, Padigaru M, Patty
Peyman JA, Rastelli L, Rieger DK, Shenoy SG, Shir
Starling G, Spytek KA, Stone DJ, Tchernev VT,
, Zerhusen BD, Zhong M;
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7 2002US-0381995P.
7 2002US-0381495P.
7 2002US-0383534P.
7 2002US-0383744P.
7 2002US-0383829P.
7 2002US-0383829P.
7 2002US-0383829P.
7 2002US-0401733P.
7 2002US-0401753P.
7 2002US-00287971.
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                                                                                 protein;
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Pred. No. 1.1e-09;
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Shimkets RA;
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Human NOVI5c protein - SEQ ID 52.

NOVX; antidiabetic; anorectic; cardiant; hypotensive; antiarteriosclerotic; virucide; antibacterial; fungicide; protozoacide; noctropic, neuroprotective; antiparkinsonian; anticonvulsant; costeopathic; antilipaemic; metabolic; diabetes; obesity; infectious; antiarthmatic; antilipaemic; metabolic; diabetes; obesity; infectious; antiarthmatic; antilipaemic; metabolic; diabetes; obesity; infectious; neurodegenerative; Alzheimer's disease; Parkinson's; epilepsy; immune; osteoarthritis; haemopoletic; inflamatory skin; asthma; dyslipidaemia; neurogenesis; cell differentiation; proliferation; haemopolesis; wound healing; angiogenesis; gene therapy; chromosome mapping; tissue typing; human; NOV.

Homo sapiens.

WC2003040330-A2.

15-MAY-2003; 2001US-033566.

05-NOV-2001; 2001US-033566.

05-NOV-2001; 2001US-033568.

05-NOV-2001; 2001US-033586.

05-NOV-2001; 2001US-033586.

05-NOV-2001; 2001US-0341477P.
17-DEC-2001; 2001US-0341490.
20-DEC-2001; 2001US-0341490.
20-DEC-2001; 2001US-0341490.
20-DEC-2001; 2001US-0341490.
21-DEC-2001; 2002US-0383544.
21-DEC-2001; 2002US-0383544.
21-DEC-2001; 2002US-0383544.
21-DEC-2001; 2002US-038354.
21-DEC-2001; 2002U
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JRA-) CURAGEN CORP.

Alsobrook JP, Alvarez E, Anderson DW, Baron M, Boldog FL;
Burgess CE, Casman SJ, Chapoval A, Dhanabal M, Edinger SR, Eisen A;
Ellerman K, Ettenberg S, Gangolli EA, Gerlach VL, Gorman L;
Grosse WM, Guo X, Hackett C, Ji W, Kekuda R, Khramtsov NV;
Lepley DM, Li L, Macdougall JR, Malyankar UM, Mazur A, Mcqueeney K;
Mezes PS, Miller CE, Millet I, Mishra VS, Padigaru M, Patturajan M;
Pena CEA, Peyman JA, Rastelli L, Rieger DK, Shenoy SG, Shimkets RA;
Smithson G, Starling G, Spytek KA, Stone DJ, Tchernev VT, Twomlow N;
Vernet CAM, Zerhusen BD, Zhong M;

WPI; 2003-441555/41. N-PSDB; ADE28674.

New isolated NOVX polypeptides and polynucleotides, useful for preventing, diagnosing or treating NOVX-associated disorders, e.g. osteoarthritis, obesity, atherosclerosis, cancer, Parkinson's disease, asthma, or infections.

Claim 1; SEQ ID NO 52; 447pp; English

The invention relates to a novel isolated NOVX polypeptide. The polypeptide of the invention demonstrates, antidiabetic, anorectic, cardiant, hypotensive, antiarteriosclerotic, virucide, antibacterial, fungicide, protozoacide, nootropic, neuroprotective, antiparkinsonian, anticonvulsant, osteopathic, antiarthritic, antiinflammatory, dermatological, antiasthmatic and antilipaemic activities. The polypeptides, nucleic acid moleules and antibodies may be useful for treating or diagnosing diseases including metabolic disorders such as diabetes and obesity, infectious diseases, anorexia, cancer,

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Matches 23
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                                                             Recombinant immunogen corresp. to dimeric form of a receptor generating antibodies able to act as receptor agonists, esp. erythropoietin receptor for treating anaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
Chimeric.
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                                       Example
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                                                                                                                                                                                                                                       07-JUN-1996;
                                                                                                                                                                                                                                                                 19-DEC-1996
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                                       Page 39-41;
                                                                                                                                                          Erickson-Miller CL
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                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                               'label= EpoR-ECD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antibody; erythropoietin receptor; EpoR; immunogen;
nionein; promoter; IgGl; Fc; anaemia; therapy.
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Pred. No. 1.1e-09;
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A fusion protein (AAW08349) encoded by plasmid mtalsEpoRFc (AAT48800) comprises the human erythropoietin receptor (EpoR) extracellular domain

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QRVEILEGRTECVLSNLRGRTRY 23

Matches Query Match Best Local (

23;

Similarity

100.0%; llarity 100.0%; Conservative 0

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Mismatches

Score 117; DB 3; Pred. No. 1.3e-09;

Length

503;

Indels

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Sequence

503 AA;

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ARAB13012
IID ARB13
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                                      protein, and to a method for screening for a candidate compound which interacts with first protein. The screening method comprises contacting the candidate molecule with a labelled modified first protein and detecting the label to identify interaction of the labelled modified first protein and candidate compound. The first protein is modified to contain a peptide, represented by sequence AAB13005. The method is used to label proteins at specific sites. The present sequence represents a Q tagged crythropoietin (EPO) receptor constructed in an example of the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      fused (via a Factor Xa cleavage sequence) to the Fc portion of human IgG1. It can be expressed e.g. in transfected Drosophila S2 cells upon induction with copper sulphate. The cells secrete EpoFFc as a dimeric molecule due to the affinity of the Fc moiety for itself. The dimeric receptor can be used as an immunogen to generate antibodies (monoclonal, polyclonal, chimeric, humanised) able to act as EcoR agonists for use in
                                                                                                                                                                                                                                                                                                         Screening for a candidate compound for use in bioassays comprises contacting the candidate molecule with a labelled modified protein detecting the label to identify interaction of the two molecules.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       erythropoietin receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Site specific label; detection; interaction screening;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-DEC-2000
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                                                                                                                                                                                                                          This invention relates to methods for the site specific modification
                                                                                                                                                                                                                                                                   Example 4; Page 26; 49pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-JAN-2000; 2000WO-US001481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB13012 standard; protein; 503
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                                                                                                                                                                                                                                                                                                       detecting the
                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-499222/44.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (SMIK ) SMITHKLINE BEECHAM CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    erythropoietin (EPO)
                                                                                                                                                                                                                                                                                                                                                                                                                                Powell DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 117;
100.0%; Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Meek TD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              <u>,</u>
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RESULT 14
AARO6512
ID AARO652
AC AARO6

RESULT 15
AAR47518
ID AAR477
XX AAR47
XX AAR47
XX 25-MA
DT 25-MA
DT 24-JU
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The sequence was deduced from DNA obtd. from a clone isolated from a commercially available human geenomic cDNA library in phage Lambda Fix (Strategene). The sequence encodes a type I trans-membrane protein with binding affinity for EPO. The gene and recombinant EPO receptor produced on expression of the DNA are used to develop reagents and systems to control and study erythropoiesis. It is believed that the EPO receptor is dys-functional in individuals with Diamond Blackfan anaemia, and may be hyperactive in polycythemia vera. See also AARO6511 (murine EPO receptor). (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GEMY )
  Erythropoietin receptor; recombinant; murine; anaemia.
                                                                                                              25-MAR-2003
24-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Erythropoietin receptor and gene - used for developing reagents systems to control and study erythropoiesis.
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04-JAN-1991
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                                                        Human EPO receptor.
                                                                                                                                                                                             AAR47518;
                                                                                                                                                                                                                                                   AAR47518 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 508 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Fig 2; 53pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dandrea A,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EPO receptor.
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23; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENETICS INST INC. WHITEHEAD INST BIOMEDICAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; ! ilarity 100.0%; ! Conservative 0;
                                                                                                              (revised)
(first entry)
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(first entry)
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                                                                                                                                                                                                                                                protein; 508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 117; DB 2;
Pred. No. 1.3e-09;
; Mismatches 0;
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Search completed: May Job time : 55 secs
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                                                                                                                                                                                            Mouse erythroleukaemia (MEL) cells were used to construct a cDNA library. The cDNA was used to transfect COS-1 cells and these were screened for radioiodinated erythropoietin (EPO) binding to isolate cDNA encoding the EPO receptor. This cDNA was used as a probe to screen a human genomic cDNA library to obtain DNA encoding the human EPO receptor. The cDNA may be used to study, treat or diagnose disorders in which the EPO receptor is dysfunctional. The EPO receptor may also be used to raise antibodied or for treating hypersensitivity to EPO or who have elevated levels of EPO. The pord. is pref. used for treating anaemias, primary proliferative polycythemia and secondary polycythemias. See also AAR47517. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GEMY )
                                                                                                                                                                       Sequence 508 AA;
                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Fig 9; 24pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                     dysfunctional.
                                                                                                                                                                                                                                                                                                                                                                                                                 Recombinant DNA encoding erythropoietin receptor - used to develop prods. for study, treatment or diagnosis of disorders in which receptor is
                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1994-025409/03.
N-PSDB; AAQ53995.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     D'andrea A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-FEB-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-1991;
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                                                        194 ORVEILEGRIECVLSNLRGRIRY 216
                                                                             1 QRVEILEGRTECVLSNLRGRTRY 23
                                                                                                               L Similarity
23; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENETICS INST INC.
CHILDRENS MEDICAL CENT.
WHITEHEAD INST BIOMEDICAL RES.
                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          89US-00306503.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "putative transmembrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "mature EPO receptor"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "signal peptide"
              6
                                                                                                             100.0%; Score 117; DB 2;
100.0%; Pred. No. 1.3e-09
tive 0; Mismatches 0
                2004, 12:45:02
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6212,

Appli Appl 12, Ap

14, Appl
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1331, App
17331, App
175, App
2743, App
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20894, Appli
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d. No. is the number of results predicted by chance to have a
re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  May 6, 2004, 12:42:22 ; Search time 23 Seconds (without alignments) 51.626 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Issued_Patents_AA:*

1: /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/2/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/2/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/2/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/2/iaa/backfiles1.pep:*
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                         US-09-028-937-11
US-08-280-864A-12
US-09-022-291-12
US-09-339-838-5
US-09-339-838-7
US-08-750-293-5
US-09-039-609-4
US-09-039-609-2
US-09-134-000C-5287
US-09-732-210-1600
US-09-732-210-1600
US-09-732-210-1600
US-09-732-210-1601
US-09-18-08-129-18-3222
US-09-18-520-316-1918-32222
US-09-18-18-5203
US-09-18-18-5203
US-09-18-752-3078-8
US-09-991-326-8
US-09-991-326-8
US-09-991-326-8
US-09-991-326-8
US-09-991-326-8
US-09-991-326-8
US-09-991-326-8
US-08-986-485-2
US-08-986-485-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                         SUMMARIES
                                                                                                                                                                      Sequence 11, Appl
Sequence 12, Appl
Sequence 5, Appli
Sequence 7, Appli
Sequence 7, Appli
Sequence 8, Appli
Sequence 4, Appli
Sequence 4, Appli
Sequence 5287, Appli
Sequence 5287, Appli
Sequence 1599, Ap
Sequence 1590, Ap
Sequence 12903, A
Sequence 2, Appli
                                                                                                                                                                                                                                                                                                                                                                                                               Description
                         Sequence
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8, Appli
8, Appli
23930, Appli
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2, Appli
5, Appli
4, Appli
                                                                                                             1601, Ap
32222, A
5203, Ap
3764, Ap
; SEQUENCE CHARACTERISTICS:

LENGTH: 23 amino acids

; TYPE: amino acid

STRANDEDNESS: unknown

; TOPOLOGY: unknown

; MOLECULE TYPE: protein

US-09-028-937-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 11, Applicati
Patent No. 6333031
GENERAL INFORMATION:
                                                                                                                                                                 FILING DATE:

CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/788,820
FILING DATE: 23-JAN-1997
PRIOR APPLICATION UNMBER: US 08/701,382
APPLICATION NUMBER: US 08/701,382
FILING DATE: 22-AUG-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/612,999
FILING DATE: 08-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Silva, Robin M.
REGISTRATION NUMBER: 38,304
REGISTRATION NUMBER: 38,304
                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA

COUNTRY: USA

ZIP: 94111-4187

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Verentarent Application Data:

CURRENT APPLICATION NUMBER: US/09/028,937
                                                                                                   TELEFAX: (415) 949-8711 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: A-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Naranda, Tatjana
TITLE OF INVENTION: Receptor Derived Peptides As Modulators
TITLE OF INVENTION: Of Receptor Activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: CA
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US-08-804-227C-14
US-09-621-976-6212
US-08-754-477A-14
US-08-754-477A-15
US-08-754-477A-15
US-08-754-177A-13
US-09-711-164-38
US-09-711-164-38
US-09-758-282B-155
US-09-758-282B-155
US-09-758-282B-243
US-08-414-034-6
US-08-414-901-6
US-08-714-901-6
US-08-714-901-6
US-08-040-741-6
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Result

Pred. No. is t score greater

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Score

Query Match

Length

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488
321
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Minimum Maximum

DB DB

geq geq

length: 0 length: 2000000000

Total number Searched: Scoring table:

BLOSUM62

Gapop 10.0 ,

Gapext 0.5

Title: Perfect score:

US-10-612-885A-1 117

QRVEILEGRTECVLSNLRGRTRY

23

Sequence:

Run on: OM protein -

protein search, using sw

model

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Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 s

summaries

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US-09-092-291-12
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US-08-280-864A-12
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Best Local Similarity
Matches 23; Conserv
                                                                                                                                                                                                         Sequence 12, Application US/09092291 Patent No. 6319499
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                         Matches
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APPLICANT: Elliott, Steven G.
TITLE OF INVENTION: Erythorpoietin Receptor
TITLE OF INVENTION: Erythorpoietin Receptor
                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Elliott,
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Winter, Robert B.
REFERENCE/DOCKET NUMBER:
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                           NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
ADDRESSEE: Amgen Inc.
STREET: One Amgen Center Di
CITY: Thousand Oaks
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 1840 CITY: Thousand Oaks
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
COMPUTER READABLE FORM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 30 amino acids
TYPE: amino acid
STRANDEDNESS: single
                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                             1 QRVEILEGRTECVLSNLRGRTRY 23
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1840 Dehavilland Drive
                                                                                                                                                                                                                                                                                                                                                                         Conservative
                               U.S.A.
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                                                                                                                                           Antibodies Which Activate 
Erythorpoietin Receptor
                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 117; DB 2; 100.0%; Pred. No. 1.8e-12; ... Mismatches 0;
                                                                                                                                                                           Steven G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 117; DB 4; 100.0%; Pred. No. 1.3e-12;
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Sequence 5, Application US/09339838
Patent No. 6361998
Patent No. 6361998
GENERAL INFORMATION:
APPLICANT: Bell, David N.
APPLICANT: Meller, Susan G.
APPLICANT: Matthews, Kathryn E.
TITLE OF INVENTION: The Efficient Culture of FILE REFERENCE: 6704-83
CURRENT FILING DATE: 1999-06-25
PRIOR APPLICATION NUMBER: US/09/339,838
CURRENT FILING DATE: 1999-01-25
PRIOR APPLICATION NUMBER: CA 2,260,332
PRIOR FILING DATE: 1998-06-25
PRIOR FILING DATE: 1998-06-25
PRIOR FILING DATE: 1998-06-25
NUMBER OF SEQ ID NOS: 11
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                                                                           RESULT 5
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Sequence 7, Application US/09339838
Patent No. 6361998
GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Winter, Robert B.
REFERENCE/DOCKET NUMBER: A
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 amino acids
CTRANTERNEES amino acids
CTRANTERNEES amino acids
                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn version 3.1 SEQ ID NO 5
                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                     LENGTH: 438
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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APPLICATION NUMBER: US,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single
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COMPUTER: I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                             23;
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                                                                                                                                                                                                    23;
                                                                                                                                                                                                                    Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 117; DB 4; 100.0%; Pred. No. 1.8e-12;
                                                                                                                                                                                                  100.0%; Score 117; DB 4; 100.0%; Pred. No. 4.3e-11; tive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of Stem Cells for the Production of Hemoglok
                                                                                                                                                                                                                                     Length 438;
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APPLICANT: Bell, David N.
APPLICANT: Mueller, Susan G.
APPLICANT: Matthews, Kathryn E.

194 QRVEILEGRTECVLSNLRGRTRY 216

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                                                                                                                         ; MOLECULE TYPE: protein US-08-776-511-2
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                                                      Matches
                                                                    Query Match
Best Local Similarity
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CURRENT FILING DATE: 1999-06-25
PRIOR APPLICATION NUMBER: CA 2,260,332
PRIOR FILING DATE: 1999-01-25
PRIOR FILING DATE: 1999-01-25
PRIOR APPLICATION NUMBER: CA 2,241,576
PRIOR FILING DATE: 1998-06-25
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Young, Peter R.
APPLICANT: Erickson-Miller, Connie
TITLE OF INVENTION: Method for Obtaining Receptor Agonist
TITLE OF INVENTION: Antibodies
                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 438
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                      NAME: Jervis, Herbert H.
REGISTRATION NUMBER: 31,171
REFERENCE/DOCKET NUMBER: SB
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                             TYPE: ami
TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 709 Swederand CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/776,511 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: SmithKline Beecham Corporation- Corporate ADDRESSEE: Patents
                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                 LENGTH:
                                                                                                                                                                                                                                                                         TELEPHONE:
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QRVEILEGRTECVLSNLRGRTRY 23
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                                                     100.0%; Score 117; DB 3; llarity 100.0%; Pred. No. 4.9e-11; Conservative 0; Mismatches 0;
                                                                                                                                                                 linear
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                                                                                                                                                                                                                                                                                                                                                                                     424
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100.0%; Pred. No. 4
tive 0; Mismatche
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                                                                                         Length 488;
                                                        Indels
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Sequence 5, Application US/08850293;
Patent No. 5843726
GENERAL INFORMATION:
APPLICANT: Lee, Jong Y.
TITLE OF INVENTION: PURIFIED HUMA:
TITLE OF INVENTION: PROTEIN FRAGM
NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-850-293-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; MOLECULE TYPE: protein US-08-850-293-5
                                                                                                                                                                                                                                               US-09-039-609-4
                                                                                                                                                                                               Sequence 4, Application US/09039609
Patent No. 6107473
                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/499,643

FILING DATE: 07-JUL-1995

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/106,815

FILING DATE: 16-AUG-1993

ATTORNEY/AGENT INFORMATION:

NAME: Ellinger, Mark S.

REGISTRATION NUMBER: 34,812

REFERENCE/DOCKET NUMBER: 07004/0

TELEPHONE: 612,335-5070
                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 612/288-9696
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/850,293
FILING DATE: 05-MAY-1997
CLASSIFICATION: 435
                                                            APPLICANT: ALBONE, EARL
APPLICANT: KIKLY, KRISTINE
TITLE OF INVENTION: A KRINGLE-RELATED CLONE,
TITLE OF INVENTION: HTHEZ47
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
STREET: P.O. Box your CITY: Valley Forge
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                                         ADDRESSEE:
                                                                                                                                                                                                                                                                                                                            194 ORVEILEGRIECVLSNLRGRIRY 216
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   amino acid
XGY: linear
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                        E: Ratner & Prestia
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SOFTWARE: FASESEO for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/039,609
FILING DATE: 16-MAR-1998
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 60/060,623
FILING DATE: 01-CCT-1997
ATTORNEY/AGENT INFORMATION:
NAME: Prestia, Paul F
REGISTRATION NUMBER: 23,031
REFERENCE/DOCKET NUMBER: GP-70275
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION: EARL
APPLICANT: KIKLY, KRISTINE
TITLE OF INVENTION: A KRINGLE-RELATED CLONE,
TITLE OF INVENTION: HTHBZ47
             FILING DATE: 16-MAR-1998
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/060,623
FILING DATE: 01-OCT-1997
ATTORNEY/AGENT INFORMATION:
NAME: Prestia, Paul F
REGISTRATION NUMBER: 23,031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 321 amino acids
                                                                                                                                                     COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/039,609
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: single TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
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REFERENCE/DOCKET NUMBER: GP-70275
                                                                                                                                                                                                                                                                                                  COUNTRY: UZIP: 19482
                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Ratner & Prestia
STREET: P.O. Box 980
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Similarity 50.0%;
10; Conservative
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Pred. No. 4.1;
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CURRENT APPLICATION NUMBER: US/09/732,210
CURRENT FILING DATE: 2000-12-07
FRIOR APPLICATION NUMBER: US 60/169,513
FRIOR FILING DATE: 1999-12-07
PRIOR APPLICATION NUMBER: US 60/169,340
PRIOR FILING DATE: 1999-12-07
NUMBER OF SEQ ID NOS: 1753
SEQ ID NO 1599
LENGTH: 67
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US-09-134-000C-5287
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APPLICANT: LYND DOLCETTE-Stamm et al
APPLICANT: LYND DOLCETTE ACID AND AMINO ACID SEQUENCES RELATING TO
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
TITLE OF INVENTION: ENTEROCOCCUS FABCALIS FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 032796-032
CURRENT APPLICATION NUMBER: US/09/134,000C
CURRENT FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/055,778
PRIOR PRILING DATE: 1997-08-15
PRIOR FILING DATE: 1997-08-15
SOUTWARE: Patentin version 3.1
SEQ ID NO 5287
LENGTH: 64
TYPE: PRT
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SEQUENCE CHARACTERISTICS:
LENGTH: 458 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                       Sequence 1599, Application US/09732210
Patent No. 6573361
GENERAL INFORMATION:
APPLICANT: Bunkers, Greg J.
APPLICANT: Liang, Jihong
APPLICANT: Mittanck, Cindy A.
APPLICANT: Seale, Jeffrey W.
APPLICANT: Wu, Yonnie S.
TITLE OF INVENTION: Anti-fungal Proteins and Methods for Their Use
FILE REFERENCE: 38-21(15036)B
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Best Local Similarity 50.0
Matches 10; Conservative
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50.0%; Pred. No. 6
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Query Match
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US-09-252-991A-32067
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                                                                                    ; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-32067
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SEQ ID NO 1600
SERGTE 67
TYPE: PRT
ORGANISM: Kluyveromyces marxianus
                                                                                                                                                                                                                                                                                                                                                                      Sequence 32067, Appli
Patent No. 6551795
GENERAL INFORMATION:
APPLICANT: Marc J.
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                                                                                                                                        SEQ ID NO 32067
LENGTH: 372
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Matches 9; Conserve
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           Matches
                                             Query Match
                                                                                                                                                                       APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER: OF SEQ. ID NOS: 33142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Wu, Yonnie S.

TITLE OF INVENTION: Anti-fungal Proteins and Methods for Their Use
FILE REFERENCE: 38-21 (15036) B

CURRENT APPLICATION NUMBER: US/09/732,210

CURRENT FILING DATE: 2000-12-07

PRIOR APPLICATION NUMBER: US 60/169,513

PRIOR PILING DATE: 1999-12-07

PRIOR PILING DATE: 1999-12-07

PRIOR PILING DATE: 1999-12-07

PRIOR FILING DATE: 1999-12-07

PRIOR FILING DATE: 1999-12-07

PRIOR FILING DATE: 1999-12-07
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APPLICANT: Liang, Jihong
APPLICANT: Mittanck, Cindy A.
APPLICANT: Seale, Jeffrey W.
APPLICANT: Wu, Yonnie S.
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ORGANISM: Kluyveromyces lactis
Local Similarity 54.5%;
hes 12; Conservative
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                           Score 45; DB Pred. No. 22;
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Pred. No. 2;
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Pred. No.
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                                               Length 372;
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SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2
LENGTH: 931
TYPE: PRT
ROANISM: Actinobacillus pleuropneumoniae
US-08-624-655A-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/489,039A CURRENT FILING DATE: 2000-01-27 PRIOR APPLICATION NUMBER: US 60/117,747 PRIOR FILING DATE: 1999-01-29 NUMBER OF SEQ ID NOS: 14342 SEQ ID NO 12903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 12903, Application US/09489039A Patent No. 6610836 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL
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                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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CURRENT APPLICATION NUMBER: US/08/624,655A
CURRENT FILING DATE: 1996-03-22
PRIOR APPLICATION NUMBER: 95 00592
PRIOR FILING DATE: 1995-03-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Gary Breton et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING
TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.2004001
                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: DABAN, MONTSERRAT
APPLICANT: MEDRANO, ANDRES
APPLICANT: ESPUNA, ENRIC
APPLICANT: QUEROL, ENRIQUE
                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS:
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TYPE: PRT
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                                                                               Local
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*

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/cgn2_6/ptodata/1/pubpaa/USO9NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/USO9NEW_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/USO9NEW_PUBCOMB.pep:*
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Copyright (c) 1993 - 2004 Compugen Ltd.
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13 US-00-016-159-5
14 US-00-016-159-5
15 US-00-316-194-144
15 US-10-316-194-35
16 US-00-316-194-35
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             Sequence 11, Appl
Sequence 30, Appl
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Sequence 50, Appl
Sequence 50, Appl
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Sequence 54, Appl
Sequence 142, Appl
Sequence 142, Appl
Sequence 143, Appl
Sequence 35, Appl
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Sequence 4, Appl
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Sequence 41, Appl
Sequence 41, Appl
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ALIGNMENTS

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CURRENT APPLICATION UNMER: US/09/991,548
CURRENT FILING DATE: 2001-11-20
PRIOR APPLICATION UNMER: 09/028,937
PRIOR FILING DATE: 1998-02-24
PRIOR APPLICATION UNMER: 08/788,820
PRIOR APPLICATION UNMER: 08/71,382
PRIOR FILING DATE: 1997-01-23
PRIOR FILING DATE: 1996-03-22
PRIOR FILING DATE: 1996-08-22
PRIOR APPLICATION NUMBER: 08/612,999
PRIOR APPLICATION NUMBER: 08/612,999
PRIOR FILING DATE: 1996-03-08
NUMBER OF SEQ ID NOS: 44
SOPTWARE: FASUSEQ for Windows Version 4.0
SEQ ID NO 11
LENGTH: 23
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                                                                                   Matches
                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: OLSSON, Lennart
APPLICANT: NARANDA, Tatjana
TITLE OF INVENTION: RECEPTOR DERIVED PEPTIDES AS MODULATORS
TITLE OF INVENTION: OF RECEPTOR ACTIVITY
FILE REFERENCE: 213542000101
                                                                                                                                                                             FEATURE: OTHER INFORMATION: human erythropoietin receptor
                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                 TYPE: PRT
                                                                                                          Similarity
                       QRVEILEGRIECVLSNLRGRIRY 23
                                                                                   Conservative
                                                                                                        100.0%; Score 117; DB 9; 100.0%; Pred. No. 1.8e-11;
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RESULT 4
US-10-287-971-48
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SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 12
LENGTH: 30
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Publication No. US20040023334A1
GENERAL INFORMATION:
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Publication No. US20030215444A1
GENERAL INFORMATION:
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 30
LENGTH: 23
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TITLE OF INVENTION: Modified Transferrin Fusion Proteins
FILE REFERENCE: 54710-5001-US
CURRENT APPLICATION NUMBER: US/10/231,494
CURRENT FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US 60/315,745
PRIOR PILING DATE: 2001-08-30
PRIOR PILING DATE: 2001-08-30
PRIOR PILING DATE: 2001-08-30
PRIOR PILING DATE: 2001-11-30
PRIOR FILING DATE: 2001-11-30
PRIOR FILING DATE: 2001-11-30
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PRIOR FILING DATE: 1998-06-05
PRIOR APPLICATION NUMBER: 08/280,864
PRIOR FILING DATE: 1994-07-26
PRIOR APPLICATION NUMBER: 09/640,090
PRIOR FILING DATE: 2000-08-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Elliott, Steven G
TITLE OF INVENTION: Antibodies which Activate an Erythropoietin Receptor
FILE REFERENCE: 06843-0030-04000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/364,276
CURRENT FILING DATE: 2003-06-17
                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: peptide
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Pred. No. 1.8e-11;
; Mismatches 0;
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RESULT 5
US-10-287-971-50
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-287-971-48
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                                                                    SOFTWARE: CuraSeqList version 0.1 SEQ ID NO 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 100.0%; Score 117; DB 12; Best Local Similarity 100.0%; Pred. No. 5.6e-10;
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APPLICANT: Alsobrook, et al
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                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/287,971
CURRENT FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 09/997,425
PRIOR FILING DATE: 2001-11-29
PRIOR APPLICATION NUMBER: 10/035,568
PRIOR FILING DATE: 2001-10-22
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PRIOR FILING DATE: 2001-11-29
PRIOR APPLICATION NUMBER: 10/035,568
PRIOR FILING DATE: 2001-10-22
                                                                                                                                              PRIOR FILING DATE: 2002-07-02
PRIOR APPLICATION NUMBER: 60/406,181
PRIOR FILING DATE: 2002-08-26
                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/348,283 PRIOR FILING DATE: 2001-11-09 PRIOR APPLICATION NUMBER: 60/393,262 PRIOR FILING DATE: 2002-07-02
                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/401,479
PRIOR FILING DATE: 2002-08-06
PRIOR APPLICATION NUMBER: 60/333,072
PRIOR FILING DATE: 2001-11-06
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CURRENT FILING DATE: 2002-11-05
                                                                                                                       NUMBER OF SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: 21402-480A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Alsobrook, et al TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NUCLEIC ACIDS ENCODING SAME, AND METHODS
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PRIOR APPLICATION NUMBER: 60/401,479
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ORGANISM: Homo sapiens
                           LENGTH: 458
TYPE: PRT
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US-10-287-971-52
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PRIOR FILLING DATE: 2001-11-29
PRIOR PELLING DATE: 2001-10-23
PRIOR FILLING DATE: 2001-10-22
PRIOR PELLING DATE: 2001-10-22
PRIOR PELLING DATE: 2001-11-05
PRIOR PELLING DATE: 2001-11-05
PRIOR PELLING DATE: 2002-08-06
PRIOR PELLING DATE: 2002-08-06
PRIOR PELLING DATE: 2001-11-06
PRIOR PELLING DATE: 2001-11-06
PRIOR PELLING DATE: 2001-11-06
PRIOR PELLING DATE: 2001-11-09
PRIOR PELLING DATE: 2002-07-02
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SOFTWARE: CuraSeqList version 0.1
SEQ ID NO 52
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                                                                                                                                                                                                                                                                                                                   Sequence 5, Application US/09016159
Patent No. US20020031806A1
GENERAL INFORMATION:
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Best Local (
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CURRENT FILING DATE: 2002-11-05
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                                                                                                                                                                                                                                                  APPLICANT: Lee, Jong Y.
TITLE OF INVENTION: PURIFIED HUMAN ERYTHROPOIETIN RECEPTOR
TITLE OF INVENTION: PROTEIN FRAGMENT AND ANTIBODIES DERIVED THEREFROM
                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                       NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C., P.A.
STREET: 60 South Sixth Street, Suite 3300
                                                                                                                          STREET: 60 South a
COMPUTER:
                                                                                        COUNTRY:
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IBM compatible
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PRIOR APPLICATION NUMBER: 09/997,425
PRIOR FILING DATE: 2001-11-29
PRIOR APPLICATION NUMBER: 10/035,568
PRIOR APPLICATION NUMBER: 60/338,626
PRIOR FILING DATE: 2001-10-22
PRIOR PILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: 60/401,479
PRIOR APPLICATION NUMBER: 60/401,479
PRIOR FILING DATE: 2002-08-06
PRIOR APPLICATION NUMBER: 60/33,072
PRIOR APPLICATION NUMBER: 60/348,283
PRIOR APPLICATION NUMBER: 60/348,283
PRIOR FILING DATE: 2001-11-09
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SOFTWARE: CuraSeqList ver
SEQ ID NO 54
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Best Local Similarity
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Query Match
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PRIOR FILING DATE: 2002-08-26
NUMBER OF COLUMBER OF C
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                                                                                                                        TYPE: PRT
ORGANISM: Homo
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LENGTH: 508 amino acids
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 612/335-5070
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FILING DATE: 21-OCT-1996
PRIOR APPLICATION NOMBER: 08/460,525
APPLICATION NUMBER: 08/460,525
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SOFTWARE: FastSE(
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               194 ORVEILEGRIECVLSNLRGRIRY 216
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No. US20040067882A1
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16-JUN-1997
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    96.6%;
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    Score 113;
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        DB 12;
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    Length 508;
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CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 10/016,516
PRIOR FILING DATE: 2001-12-10
NUMBER OF SEQ ID NOS: 173
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 142
                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 173
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 144
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 144, Application US/10316194 Publication No. US20030215914A1
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Best Local
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                                                                                                                                                                                                                                                                                                                                        APPLICANT: Houtzager, Erwin
APPLICANT: Vijn, Irma M.C.
APPLICANT: Sijmons, Peter C.
TITLE OF INVENTION: A structure for presenting desired peptide sequences
FILE REFERENCE: 2183-5610US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Houtzager, Erwin
APPLICANT: Vijn, Irma M.C.
APPLICANT: Sijmons, Peter C.
TITLE OF INVENTION: A structure for presenting desired peptide sequences
FILE REFERENCE: 2183-5610US
                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/316,194
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 10/016,516
PRIOR FILING DATE: 2001-12-10
                                                                      FEATURE:
NAME/KEY: SITE
LOCATION: (1)...
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NAME/KEY: SITE
LOCATION: (1)..
                                                                                                                                                         TYPE: PRT ORGANISM: Artificial Sequence
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                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence: IMABIS027
                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: IMABIS020
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                                                                                                                                                                                                   ENGTH: 109
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b. US20030215914A1
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82.1%;
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1; Mismatches 0
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Pred. No. 1.8e-08;
Score 96; DB 15;
Pred. No. 2.3e-07;
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                 Length 109;
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; Sequence 35, Application US/10316194
"""h14-ation No. US20030215914A1
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CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 10/016,516
PRIOR FILING DATE: 2001-12-10
NUMBER OF SEQ ID NOS: 173
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 43
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Publication No. US20030215914A1
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SEQ ID NO 35
LENGTH: 109
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Query Match
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APPLICANT: Vijn, Irma M.C.
APPLICANT: Sijmons, Peter C.
TITLE OF INVENTION: A structure for presenting desired peptide sequences
FILE REFERENCE: 2183-5610US
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CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 10/016,516
PRIOR FILING DATE: 2001-12-10
NUMBER OF SEQ ID NOS: 173
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APPLICANT: Sijmons, Peter C.
TITLE OF INVENTION: A structure for presenting desired peptide sequences
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                                                                                   OTHER INFORMATION: Description of Artificial Sequence: VAP amino acid OTHER INFORMATION: sequence of iMab1100
                                                                                                                                                       TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: VAP amino acid
OTHER INFORMATION: sequence of iMab600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT ORGANISM: Artificial Sequence
                                                   LOCATION:
                                                                    NAME/KEY: SITE
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NAME/KEY: SITE
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 73.5%;
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 Score 86;
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Pred. No.
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Length 109;
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SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 11
LENGTH: 1026
TYPE: PRT
ORGANISM: Homo sapiens
US-09-947-063-11
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RESULT 15
US-09-728-912-7
; Sequence 7, Application US/09728912
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-947-063-4
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CURRENT APPLICATION NUMBER: US/09/947,063
CURRENT FILING DATE: 2001-09-05
PRIOR APPLICATION NUMBER: 60/229,990
PRIOR FILING DATE: 2000-09-05
PRIOR APPLICATION NUMBER: 60/229,988
PRIOR FILING DATE: 2000-09-05
PRIOR APPLICATION NUMBER: 60/229,988
PRIOR FILING DATE: 2000-09-05
PRIOR PLING DATE: 2001-09-05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 11, Application US/09947063
Publication No. US20030059775A1
GENERAL INFORMATION:
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Best Local Similarity 71.4%;
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TITLE OF INVENTION: No. US20030059775A1el Proteins and Nucleic Acids Encoding Same
FILE REFERENCE: 21402-112
CURRENT APPLICATION NUMBER: US/09/947,063
CURRENT FILING DATE: 2001-09-05
PRIOR APPLICATION NUMBER: 60/229,990
PRIOR APPLICATION NUMBER: 60/229,988
PRIOR PILING DATE: 2000-09-05
PRIOR PILING DATE: 2000-09-05
PRIOR FILING DATE: 2000-09-05
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TITLE OF INVENTION: No. US20030059775Alel Proteins and Nucleic Acids Encoding Same
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 30
                                                                                                                                                                                          292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   292 EGFYECIASNIRGR 305
                                                                                                                                                                                                                                                                                                                                         10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10;
                                                                                                                                                                                                                                               7 EGRTECVLSNLRGR 20
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                                                                                                                                                                                          EGFYECIASNLRGR 305
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                                                                                                                                                                                                                                                                                                                                 Score 52; DB 10;
pred. No. 27;
1; Mismatches 3
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Pred. No. 27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred. No. 8.6e-06; 0; Mismatches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                                         Indels
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APPLICANT: HOLLOWAY, James L.
APPLICANT: HOLLOWAY, James L.
TITLE OF INVENTION: Tumor Necrosis Factor-Stimulated Gene
TITLE OF INVENTION: and Protein
FILE REFERENCE: 99-94US
CURRENT APPLICATION NUMBER: US/09/728,912
CURRENT FILING DATE: 2000-12-01
PRIOR APPLICATION NUMBER: 60/169,252
PRIOR FILING DATE: 1999-12-06
NUMBER OF SEQ ID NOS: 14
SOFTMARE: FastSEQ for Windows Version 3.0
SEQ ID NO 7
LENGTH: 49
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                                                                                                                                         US-09-728-912-7
                                                                                                                                                                                                                                                                                                                                                                                                      Patent No. US20010036643A1
GENERAL INFORMATION:
                                                                                    Query Match
Best Local Similarity
                                                                     Matches
                                                                                                                                                        LENGTH: 49
TYPE: PRT
ORGANISM: Homo sapiens
                                                                     10;
                  3 VEILEGRIECVLSNLRGRIR 22
7 VELLDGYTHRVLARFHGRSR 26
                                                                     Conservative
                                                                                    41.9%;
                                                                     4; Mismatches
                                                                                    Score 49; I
Pred. No. 2.
                                                                                                      DB 9;
                                                                     6; Indels
                                                                                                      Length 49;
                                                                     0
                                                                     Gaps
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Search completed: May 6, 2004, 12:48:22 Job time: 43 secs

